```
   1 CCATGGGAGC GAACACTTCA AGAAAACCAC CAGTGTTTGA TGAAAATGAA
  51 GATGTCAACT TTGACCACTT TGAAATTTTG CGAGCCATTG GGAAAGGCAG
 101 TTTTGGGGAG GTCTGCATTG TACAGAAGAA TGATACCAAG AAGATGTGCG
 151 CAATGAAGTA CATGAATAAA CAAAAGTGCG TGGAGCGCAA TGAAGTGAGA
 201 AATGTCTTCA AGGAACTCCA GATCATGCAG GGTCTGGAGC ACCCTTTCCT
 251 GGTTAATTTG TGGTATTCCT TCCAAGATGA GGAAGACATG TTCATGGTGG
 301 TGGACCTCCT GCTGGGTGGA GACCTGCGTT ATCACCTGCA ACAGAACGTC
 351 CACTTCAAGG AAGAAACAGT GAAGCTCTTC ATCTGTGAGC TGGTCATGGC
 401 CCTGGACTAC CTGCAGAACC AGCGCATCAT TCACAGGGAT ATGAAGCCTG
 451 ACAATATTTT ACTTGACGAA CATGGGCACG TGCACATCAC AGATTTCAAC
 501 ATTGCTGCGA TGCTGCCCAG GGAGACACAG ATTACCACCA TGGCTGGCAC
 551 CAAGCCTTAC ATGGCACCTG AGATGTTCAG CTCCAGAAAA GGAGCAGGCT
 601 ATTCCTTTGC TGTTGACTGG TGGTCCCTGG GAGTGACGGC ATATGAACTG
 651 CTGAGAGGCC GGAGACCGTA TCATATTCGC TCCAGTACTT CCAGCAAGGA
 701 AATTGTACAC ACGTTTGAGA CGACTGTTGT AACTTACCCT TCTGCCTGGT
 751 CACAGGAAAT GGTGTCACTT CTTAAAAAGC TACTCGAACC TAATCCAGAC
 801 CAACGATTTT CTCAGTTATC TGATGTCCAG AACTTCCCGT ATATGAATGA
 851 TATAAACTGG GATGCAGTTT TTCAGAAGAG GCTCATTCCA GGTTTCATTC
 901 CTAATAAAGG CAGGCTGAAT TGTGATCCTA CCTTTGAACT TGAGGAAATG
 951 ATTTTGGAGT CCAAACCTCT ACATAAGAAA AAAAAGCGTC TGGCAAAGAA
1001 GGAGAAGGAT ATGAGGAAAT GCGATTCTTC TCAGACATGT CTTCTTCAAG
1051 AGCACCTTGA CTCTGTCCAG AAGGAGTTCA TAATTTTCAA CAGAGAAAAA
1101 GTAAACAGGG ACTTTAACAA AAGACAACCA AATCTAGCCT TGGAACAAAC
1151 CAAAGACCCA CAAGGTGAGG ATGGTCAGAA TAACAACTTG TAAAGGCCTC
1201 ATGTCTTCTT CTTGGGACAA TCTCATGCCA GAAACTTCTA ATTACATATG
1251 TCAAGAAAAG CTGACAGTAG CTCCTGCCAC TCCACACACC ATGACTTAGA
1301 AAATGTGAAT GAATATATTT CAAAAAAGGC AGCACAACAC AGTGAAGGGT
1351 CCTGGGCCTG AGCTCCTGGA AAGTCATTTC ACATCAATCA ACTGTGTGAT
1401 CTAGAGCAAG TCACTTAGCC ACTTTCTGTG CTTTACTTTA TTTATCTAAA
1451 ATGAGAGGGT TATACTAAAA AAAAAAAAAA AAAAA
```
   (SED NO:1)
5'UTR: 1 - 2
Start Codon: 3
Stop Codon: 1191
3'UTR: 1194

Homologous proteins:

Top 10 BLAST Hits	Score	E
CRA|87000000001426 /altid=gi|7161864 /def=emb|CAB76566.1| (AJ25...	560	e-158
CRA|87000000001314 /altid=gi|8923754 /def=ref|NP_060871.1| gene...	557	e-157
CRA|103000001515936 /altid=gi|10946600 /def=ref|NP_067277.1| hy...	514	e-145
CRA|108000024647823 /altid=gi|12730486 /def=ref|XP_003392.2| ge...	395	e-109
CRA|18000005184360 /altid=gi|7505957 /def=pir||T23688 hypotheti...	328	8e-89
CRA|18000005004115 /altid=gi|1730069 /def=sp|P54644|KRAC_DICDI ...	226	5e-58
CRA|18000004912236 /altid=gi|464395 /def=sp|P28178|PK2_DICDI PR...	209	8e-53
CRA|18000004991065 /altid=gi|1362152 /def=pir||S56639 ribosomal...	204	3e-51
CRA|18000004952305 /altid=gi|462434 /def=sp|P34099|KAPC_DICDI C...	203	6e-51
CRA|107000045076305 /altid=gi|12322721 /def=gb|AAG51345.1|AC012...	202	8e-51

EST:	Score	E
gi|12432521 /dataset=dbest /taxon=96...	1362	0.0
gi|12425892 /dataset=dbest /taxon=96...	864	0.0
gi|9811536 /dataset=dbest /taxon=960...	708	0.0

EXPRESSION INFORMATION FOR MODULATORY USE:
gi\|12432521 brain hippocampus
gi\|12425892 Breast mammary adenocarcinoma cell line
gi\|9811536 Bladder carcinoma cell line
Tissue expression:
Human brain
Human fetal brain
Human fetal heart
Human kidney
Human uterus

FIGURE 1

```
  1 MGANTSRKPP VFDENEDVNF DHFEILRAIG KGSFGEVCIV QKNDTKKMCA
 51 MKYMNKQKCV ERNEVRNVFK ELQIMQGLEH PFLVNLWYSF QDEEDMFMVV
101 DLLLGGDLRY HLQQNVHFKE ETVKLFICEL VMALDYLQNQ RIIHRDMKPD
151 NILLDEHGHV HITDFNIAAM LPRETQITTM AGTKPYMAPE MFSSRKGAGY
201 SFAVDWWSLG VTAYELLRGR RPYHIRSSTS SKEIVHTFET TVVTYPSAWS
251 QEMVSLLKKL LEPNPDQRFS QLSDVQNFPY MNDINWDAVF QKRLIPGFIP
301 NKGRLNCDPT FELEEMILES KPLHKKKKRL AKKEKDMRKC DSSQTCLLQE
351 HLDSVQKEFI IFNREKVNRD FNKRQPNLAL EQTKDPQGED GQNNNL
   (SEQ ID NO:2)
```

FEATURES:
Functional domains and key regions:
[1] PDOC00001 PS00001 ASN_GLYCOSYLATION
N-glycosylation site

Number of matches: 2
```
        1          4-7 NTSR
        2        43-46 NDTK
```
----
[2] PDOC00005 PS00005 PKC_PHOSPHO_SITE
Protein kinase C phosphorylation site

Number of matches: 7
```
        1          5-7 TSR
        2          6-8 SRK
        3      194-196 SRK
        4        45-47 TKK
        5      122-124 TVK
        6      193-195 SSR
        7          6-8 SRK
```
----
[3] PDOC00006 PS00006 CK2_PHOSPHO_SITE
Casein kinase II phosphorylation site

Number of matches: 4
```
        1        33-36 SFGE
        2        89-92 SFQD
        3      212-215 TAYE
        4      230-233 SSKE
```
----
[4] PDOC00008 PS00008 MYRISTYL
N-myristoylation site

Number of matches: 3
```
        1          2-7 GANTSR
        2      197-202 GAGYSF
        3      391-396 GQNNNL
```
----
[5] PDOC00009 PS00009 AMIDATION
Amidation site

```
          218-221 RGRR
```
----
[6] PDOC00100 PS00107 PROTEIN_KINASE_ATP
Protein kinases ATP-binding region signature

```
          29-52 IGKGSFGEVCIVQKNDTKKMCAMK
```
----
[7] PDOC00100 PS00108 PROTEIN_KINASE_ST
Serine/Threonine protein kinases active-site signature

```
          142-154 IIHRDMKPDNILL
```

Membrane spanning structure and domains:
```
   Helix Begin    End    Score Certainity
     1    197     217    0.690 Putative
```

FIGURE 2A

BLAST Alignment t T p Hit:
Alignment to top blast hit:
>CRA|87000000001426 /altid=gi|7161864 /def=emb|CAB76566.1| (AJ250840)
          serine/threonine protein kinase [Mus musculus] /org=Mus
          musculus /taxon=10090 /dataset=nraa /length=414
            Length = 414

 Score = 560 bits (1428), Expect = e-158
 Identities = 278/403 (68%), Positives = 320/403 (78%), Gaps = 7/403 (1%)
 Frame = +3

Query: 3   MGANTSRKPPVFDENEDVNFDHFEILRAIGKGSFGEVCIVQKNDTKKMCAMKYMNKQKCV 182
             MG N S KPPVFDENE+VNFDHF+ILRAIGKGSFG+VCIVQK DTKKM AMKYMNKQKCV
Sbjct: 1   MGGNHSHKPPVFDENEEVNFDHFQILRAIGKGSFGKVCIVQKRDTKKMYAMKYMNKQKCV 60

Query: 183   ERNEVRNVFKELQIMQGLEHPFLVNLWYSFQDEEDMFMVVDLLLGGDLRYHLQQNVHFKE 362
             ER+EVRNVF+ELQIMQGLEHPFLVNLWYSFQDEEDMFMVVDLLLGGDLRYHLQQNVHF E
Sbjct: 61   ERDEVRNVFRELQIMQGLEHPFLVNLWYSFQDEEDMFMVVDLLLGGDLRYHLQQNVHFTE 120

Query: 363   ETVKLFICELVMALDYLQNQRIIHRDMKPDNILLDEHGHVHITDFNIAAMLPRETQITTM 542
             TVKL+ICEL +AL+YLQ   IIHRD+KPDNILLDEHGHVHITDFNIA +L   + ++M
Sbjct: 121   GTVKLYICELALALEYLQRYHIIHRDIKPDNILLDEHGHVHITDFNIATVLKGSEKASSM 180

Query: 543   AGTKPYMAPEMFS--SRKGAGYSFAVDWWSLGVTAYELLRGRRPYHIRSSTSSKEIVHTF 716
             AGTKPYMAPE+F   G GYS+ VDWWSLGVTAYELLRG RPY I S+T   EI++ F
Sbjct: 181   AGTKPYMAPEVFQVYVDGGPGYSYPVDWWSLGVTAYELLRGWRPYEIHSATPIDEILNMF 240

Query: 717   ETTVVTYPSAWSQEMVSLLKKLLEPNPDQRFSQLSDVQNFPYMNDINWDAVFQKRLIPGF 896
             + V Y S W + MVSLLKKLL +P+ R S L D+Q+ Y+ D+NWDAVF+K L+PGF
Sbjct: 241   KVERVHYSSTWCEGMVSLLKKLLTKDPESRLSSLRDIQSMTYLADMNWDAVFEKALMPGF 300

Query: 897   IPNKGRLNCDPTFELEEMILESKPLHKKKKRLAK-KEKDMRKCDSSQTCLLQEHLDSVQK 1073
             +PNKGRLNCDPTFELEEMILESKPLHKKKKRLAK + +D K   LQ+ L++V+K
Sbjct: 301   VPNKGRLNCDPTFELEEMILESKPLHKKKKRLAKHRSRDSTKDSCPLNGHLQQCLETVRK 360

Query: 1074 EFIIFNREKVNR----DFNKRQPNLALEQTKDPQGEDGQNNNL 1190
             EFIIFNREK+ R   D   + +   + +DG+NNN+
Sbjct: 361   EFIIFNREKLRRQQGHDGQLSDLDGRIGSQTSSKLQDGRNNNI 403 (SEQ ID NO:4)

>CRA|87000000001314 /altid=gi|8923754 /def=ref|NP_060871.1| gene for
          serine/threonine protein kinase [Homo sapiens] /org=Homo
          sapiens /taxon=9606 /dataset=nraa /length=414
            Length = 414

 Score = 557 bits (1419), Expect = e-157
 Identities = 275/403 (68%), Positives = 319/403 (78%), Gaps = 7/403 (1%)
 Frame = +3

Query: 3   MGANTSRKPPVFDENEDVNFDHFEILRAIGKGSFGEVCIVQKNDTKKMCAMKYMNKQKCV 182
             MG N S KPPVFDENE+VNFDHF+ILRAIGKGSFG+VCIVQK DTKKM AMKYMNKQKC+
Sbjct: 1   MGGNHSHKPPVFDENEEVNFDHFQILRAIGKGSFGKVCIVQKRDTKKMYAMKYMNKQKCI 60

Query: 183   ERNEVRNVFKELQIMQGLEHPFLVNLWYSFQDEEDMFMVVDLLLGGDLRYHLQQNVHFKE 362
             ER+EVRNVF+ELQIMQGLEHPFLVNLWYSFQDEEDMFMVVDLLLGGDLRYHLQQNVHF E
Sbjct: 61   ERDEVRNVFRELQIMQGLEHPFLVNLWYSFQDEEDMFMVVDLLLGGDLRYHLQQNVHFTE 120

Query: 363   ETVKLFICELVMALDYLQNQRIIHRDMKPDNILLDEHGHVHITDFNIAAMLPRETQITTM 542
             TVKL+ICEL +AL+YLQ   IIHRD+KPDNILLDEHGHVHITDFNIA ++   + ++M
Sbjct: 121   GTVKLYICELALALEYLQRYHIIHRDIKPDNILLDEHGHVHITDFNIATVVKGAERASSM 180

Query: 543   AGTKPYMAPEMFS--SRKGAGYSFAVDWWSLGVTAYELLRGRRPYHIRSSTSSKEIVHTF 716
             AGTKPYMAPE+F   +G GYS+ VDWWSLG+TAYELLRG RPY I S T   EI++ F
Sbjct: 181   AGTKPYMAPEVFQVYMDRGPGYSYPVDWWSLGITAYELLRGWRPYEIHSVTPIDEILNMF 240

Query: 717   ETTVVTYPSAWSQEMVSLLKKLLEPNPDQRFSQLSDVQNFPYMNDINWDAVFQKRLIPGF 896
             + V Y S W + MV+LL+KLL +P+ R S L D+Q+ PY+ D+NWDAVF+K L+PGF
Sbjct: 241   KVERVHYSSTWCKGMVALLRKLLTKDPESRVSSLHDIQSVPYLADMNWDAVFKKALMPGF 300

Query: 897   IPNKGRLNCDPTFELEEMILESKPLHKKKKRLAK-KEKDMRKCDSSQTCLLQEHLDSVQK 1073
             +PNKGRLNCDPTFELEEMILESKPLHKKKKRLAK + +D K   LQ L++V++
Sbjct: 301   VPNKGRLNCDPTFELEEMILESKPLHKKKKRLAKNRSRDGTKDSCPLNGHLQHCLETVRE 360

Query: 1074 EFIIFNREKVNRDFNKRQPNLALEQTKDPQG----EDGQNNNL 1190
             EFIIFNREK+ R + L +   Q   +DG NNNL
Sbjct: 361   EFIIFNREKLRRQQGQGSQLLDTDSRGGGQAQSKLQDGCNNNL 403 (SEQ ID NO:5)

FIGURE 2B

>CRA|103000001515936 /altid=gi|10946600 /def=ref|NP_067277.1|
           hypothetical serine/threonine protein kinase [Mus
           musculus] /org=Mus musculus /taxon=10090 /dataset=nraa
           /length=488
           Length = 488

   Score = 514 bits (1310), Expect = e-145
   Identities = 250/389 (64%), Positives = 304/389 (77%), Gaps = 4/389 (1%)
   Frame = +3

Query: 18   SRKPPVFDENEDVNFDHFEILRAIGKGSFGEVCIVQKNDTKKMCAMKYMNKQKCVERNEV 197
             S + PVFD+ EDVNFDHF+ILRAIGKGSFG+VCIVQK DT+KM AMKYMNKQ+C+ER+EV
Sbjct: 77   SARRPVFDDKEDVNFDHFQILRAIGKGSFGKVCIVQKRDTEKMYAMKYMNKQQCIERDEV 136

Query: 198   RNVFKELQIMQGLEHPFLVNLWYSFQDEEDMFMVVDLLLGGDLRYHLQQNVHFKEETVKL 377
             RNVF+EL+I+Q +EH FLVNLWYSFQDEEDMFMVVDLLLGGDLRYHLQQNV F E+TV+L
Sbjct: 137   RNVFRELEILQEIEHVFLVNLWYSFQDEEDMFMVVDLLLGGDLRYHLQQNVQFSEDTVRL 196

Query: 378   FICELVMALDYLQNQRIIHRDMKPDNILLDEHGHVHITDFNIAAMLPRETQITTMAGTKP 557
             +ICE+ +ALDYL++Q IIHRD+KPDNILLDE GH H+TDFNIA ++   + T +AGTKP
Sbjct: 197   YICEMALALDYLRSQHIIHRDVKPDNILLDEQGHAHLTDFNIATIIKDGERATALAGTKP 256

Query: 558   YMAPEMFSS--RKGAGYSFAVDWWSLGVTAYELLRGRRPYHIRSSTSSKEIVHTFETTVV 731
             YMAPE+F S   G GYSF VDWWS+GV AYELLRG RPY I SS + + +V F T V
Sbjct: 257   YMAPEIFHSFVNGGTGYSFEVDWWSVGVMAYELLRGWRPYDIHSSNAVESLVQLFSTVSV 316

Query: 732   TYPSAWSQEMVSLLKKLLEPNPDQRFSQLSDVQNFPYMNDINWDAVFQKRLIPGFIPNKG 911
               Y   WS+EMV+LL+KLL NP+ RFS L D+Q P + + WD + +K++ PGF+PNKG
Sbjct: 317   QYVPTWSKEMVALLRKLLTVNPEHRFSSLQDMQTAPSLAHVLWDDLSEKKVEPGFVPNKG 376

Query: 912   RLNCDPTFELEEMILESKPLHKKKKRLAKKEKDMRKCDSSQT--CLLQEHLDSVQKEFII 1085
             RL+CDPTFELEEMILES+PLHKKKKRLAK +   DSSQ+   LQ+ LD++Q++F+I
Sbjct: 377   RLHCDPTFELEEMILESRPLHKKKKRLAKNKSRDSSRDSSQSENDYLQDCLDAIQQDFVI 436

Query: 1086 FNREKVNRDFNKRQPNLALEQTKDPQGED 1172
             FNREK+   KR L E   P+ D
Sbjct: 437   FNREKL-----KRSQELMSEPPPGPETSD 460 (SEQ ID NO:6)

>CRA|108000024647823 /altid=gi|12730486 /def=ref|XP_003392.2| gene for
           serine/threonine protein kinase [Homo sapiens] /org=Homo
           sapiens /taxon=9606 /dataset=nraa /length=330
           Length = 330

   Score = 395 bits (1004), Expect = e-109
   Identities = 199/316 (62%), Positives = 237/316 (74%), Gaps = 7/316 (2%)
   Frame = +3

Query: 264   YSFQDEEDMFMVVDLLLGGDLRYHLQQNVHFKEETVKLFICELVMALDYLQNQRIIHRDM 443
             YSFQDEEDMFMVVDLLLGGDLRYHLQQNVHF E TVKL+ICEL +AL+YLQ   IIHRD+
Sbjct: 4   YSFQDEEDMFMVVDLLLGGDLRYHLQQNVHFTEGTVKLYICELALALEYLQRYHIIHRDI 63

Query: 444   KPDNILLDEHGHVHITDFNIAAMLPRETQITTMAGTKPYMAPEMFS--SRKGAGYSFAVD 617
             KPDNILLDEHGHVHITDFNIA ++   + ++MAGTKPYMAPE+F   +G GYS+ VD
Sbjct: 64   KPDNILLDEHGHVHITDFNIATVVKGAERASSMAGTKPYMAPEVFQVYMDRGPGYSYPVD 123

Query: 618   WWSLGVTAYELLRGRRPYHIRSSTSSKEIVHTFETTVVTYPSAWSQEMVSLLKKLLEPNP 797
             WWSLG+TAYELLRG RPY I S T   EI++ F+   V Y S W + MV+LL+KLL +P
Sbjct: 124   WWSLGITAYELLRGWRPYEIHSVTPIDEILNMFKVERVHYSSTWCKGMVALLRKLLTKDP 183

Query: 798   DQRFSQLSDVQNFPYMNDINWDAVFQKRLIPGFIPNKGRLNCDPTFELEEMILESKPLHK 977
             + R S L D+Q+ PY+ D+NWDAVF+K L+PGF+PNKGRLNCDPTFELEEMILESKPLHK
Sbjct: 184   ESRVSSLHDIQSVPYLADMNWDAVFKKALMPGFVPNKGRLNCDPTFELEEMILESKPLHK 243

Query: 978   KKKRLAK-KEKDMRKCDSSQTCLLQEHLDSVQKEFIIFNREKVNRDFNKRQPNLALEQTK 1154
             KKKRLAK + +D K   LQ L++V++EFIIFNREK+ R + L +
Sbjct: 244   KKKRLAKNRSRDGTKDSCPLNGHLQHCLETVREEFIIFNREKLRRQQGQGSQLLDTDSRG 303

Query: 1155 DPQG----EDGQNNNL 1190
             Q   +DG NNNL
Sbjct: 304   GGQAQSKLQDGCNNNL 319 (SEQ ID NO: 7)

>CRA|18000005184360 /altid=gi|7505957 /def=pir||T23688 hypothetical
           protein M03C11.1 - Caenorhabditis elegans
           /org=Caenorhabditis elegans /taxon=6239 /dataset=nraa
           /length=379
           Length = 379

FIGURE 2C

```
Score =  328 bits (833), Expect = 8e-89
Identities = 156/353 (44%), Positives = 226/353 (63%), Gaps = 2/353 (0%)
Frame = +3

Query: 66    HFEILRAIGKGSFGEVCIVQKNDTKKMCAMKYMNKQKCVERNEVRNVFKELQIMQGLEHP 245
             HF ++R+IG+G+FG+VCIVQ+  TKK  A+KYMNK++C+E+    NV +EL ++  + HP
Sbjct: 27    HFSVIRSIGRGAFGKVCIVQERKTKKYFALKYMNKRRCIEKGVAANVIRELTLLSKMSHP 86

Query: 246   FLVNLWYSFQDEEDMFMVVDLLLGGDLRYHLQQNVHFKEETVKLFICELVMALDYLQNQR 425
             F+VNLWY+FQD + M+MV DLLLGGDLRYHL Q  F E+  KL++CE+ +A++YL   +
Sbjct: 87    FIVNLWYTFQDGDYMYMVSDLLLGGDLRYHLSQQGKFAEDRAKLYLCEICLAVEYLHEMK 146

Query: 426   IIHRDMKPDNILLDEHGHVHITDFNIAAMLPRETQITTMAGTKPYMAPEMFSS--RKGAG 599
             I+HRD+KP+NILLDE GH H+TD N+A  L  +   T+ +GT+PYMAPE++++     G
Sbjct: 147   IVHRDIKPENILLDEQGHAHLTDLNLATQLEDDQLATSYSGTRPYMAPEIYATYLEIEDG 206

Query: 600   YSFAVDWWSLGVTAYELLRGRRPYHIRSSTSSKEIVHTFETTVVTYPSAWSQEMVSLLKK 779
             Y   VDWW+LGV  YE+LRGR P+   S T  +E   F + YP+ W  +++  +
Sbjct: 207   YDSRVDWWALGVCFYEMLRGRTPFEFSSRTKPEEAYVAFRESSIPYPAHWPTDLIQFINS 266

Query: 780   LLEPNPDQRFSQLSDVQNFPYMNDINWDAVFQKRLIPGFIPNKGRLNCDPTFELEEMILE 959
             +L+ + ++R   L ++    Y   I++ +VF+K+  P FIP K  LNCDP +ELEE IL
Sbjct: 267   MLKFDKEKRLVGLEAIKKHSYTERIDFKSVFEKKPSPVFIPCKEGLNCDPMYELEERILV 326

Query: 960   SKPLHKKKKRLAKKEKDMRKCDSSQTCLLQEHLDSVQKEFIIFNREKVNRDFN 1118
             S P+H  ++R        R   Q  L E   V K FI F+R V + N
Sbjct: 327   STPIH--RRRTNHNNSSGRSSSEPQNAALVE----VSKAFIDFSRHNVKIEPN 373 (SEQ ID NO: 8)
```

<u>Hmmer search results (Pfam):</u>
Scores for sequence family classification (score includes all domains):

Model	Description	Score	E-value	N
PF00069	Eukaryotic protein kinase domain	241.7	1e-68	1
CE00359	E00359 bone_morphogenetic_protein_receptor	18.9	0.00012	1
PF00433	Protein kinase C terminal domain	8.8	0.2	1
CE00022	CE00022 MAGUK_subfamily_d	8.6	0.02	1
CE00203	CE00203 ERBB_RECEPTOR	6.0	0.25	1
CE00031	CE00031 VEGFR	5.1	0.12	1
CE00528	CE00528 CDC14_PHOSPHATASE	1.7	8	1
CE00292	CE00292 PTK_membrane_span	-44.9	1.9e-06	1
CE00287	CE00287 PTK_Eph_orphan_receptor	-45.1	2.4e-05	1
CE00286	E00286 PTK_EGF_receptor	-59.9	1.7e-07	1
CE00291	CE00291 PTK_fgf_receptor	-81.9	0.00049	1
CE00290	CE00290 PTK_Trk_family	-158.8	0.00022	1
CE00016	CE00016 GSK_glycogen_synthase_kinase	-216.0	0.00011	1
CE00288	CE00288 PTK_Insulin_receptor	-225.8	0.21	1

FIGURE 2D

Parsed for domains:

Model	Domain	seq-f	seq-t		hmm-f	hmm-t		score	E-value
CE00031	1/1	134	168	..	1059	1093	..	5.1	0.12
CE00203	1/1	136	172	..	855	891	..	6.0	0.25
CE00359	1/1	142	191	..	272	326	..	18.9	0.00012
CE00022	1/1	133	223	..	133	226	..	8.6	0.02
CE00288	1/1	23	238	..	1	269	[]	-225.8	0.21
CE00528	1/1	251	260	..	608	617	.]	1.7	8
CE00292	1/1	23	276	..	1	288	[]	-44.9	1.9e-06
CE00287	1/1	23	276	..	1	260	[]	-45.1	2.4e-05
CE00291	1/1	23	278	..	1	285	[]	-81.9	0.00049
CE00286	1/1	23	278	..	1	263	[]	-59.9	1.7e-07
CE00290	1/1	23	279	..	1	282	[]	-158.8	0.00022
PF00069	1/1	23	281	..	1	278	[]	241.7	1e-68
PF00433	1/1	282	301	..	1	20	[.	8.8	0.2
CE00016	1/1	1	331	[.	1	433	[]	-216.0	0.00011

FIGURE 2E

```
   1 TCCCTCTCTC ATACCATTTA ATTGGTTGCT TCCTAATTAA TGACTCTCTT
  51 TGCTCTCTAT TTAATGATTC TTGCTAAAGT CCATAAGGCA CTTTGCCAGC
 101 AGTTGGTTTT TAGTATGAAA AGTAGCATTT CCTTAATGAG TCTGAGTCTG
 151 CCTTCCAAAT GAAGGGTTTA CTTACATTTT CCTAATGGGA AAACGAGCTT
 201 TTCTTCTACG CTTCCTTAGG GGTTTCATAA GTTCTTTTTC AATAACTCAT
 251 CCTTAACACT TTCTCCAATT CTGCCTGTAA TCAATATTCC CTTCACATGT
 301 AAAGAGCTCA GGAGGAAATC AACTATTTTT TTAAAAATAC GCAATAAGGA
 351 AATTCTGCTA CTCTTAGAAA TAGCAGGAGC TAACATTCAT TCTTTGCATA
 401 TCATGTGCTA GGCATTGTGC CAATTACCTT ATATACATTG TCTCATTATA
 451 TGTATCCATG ACCATATATG TGCTAAGCAT GAAATTTTCT TAAGCCAGAT
 501 AGCTGAGTAG AATTTTAAAA TATTATTTTG TACAAAATCT AGACCTTTAC
 551 CCCATTTGGG GGATAGATCT GAAGATCTGG GCTCATGTTT CCATGTGGTG
 601 ACAATCTGTT TGATCTGAGC ACAATTACTT TATTTGGATG GAGCCATTGC
 651 CACCATTGTC TGCCCAATGC ACTAATGTTA AATGCCCAGT CTGGCTCACT
 701 CATTTGCATC ATCTGCCTGG CTCCTATAGG GATCCCAGCT TGTCACTCCT
 751 GAGGTAGACA CTGTCATTTC CCCCATTCTA GAGGTGAGAG GTTACATAAC
 801 TGGGCCAAAG GCATTATCAG TGTCAGTTTT AGGACTGGAA CACAGGATGC
 851 TGCCTCTCTT TACCATTATG TTTTAAAGTG GAGCAAAGCC GTAGTTTTCA
 901 GGATCTTTTC TTGTTCACAC ATATCATTTA ATTTGAGCCT CAGAGCGGCT
 951 AACAGTTTTG AGCACTTATG CTATGAAAAT GTTTTGTGTA TTCAGTTAAA
1001 TGTATGCATA TCATACATTT ATGTAACTCA ATACATATAT ATAAATGTGA
1051 TATAACATAC GTATGATATA ACAGAGTTAT ATATATGTGT ATTATTTAAC
1101 TTAATATATA ATGAGTTAAG TGTATGCATA TCATAGATTT ATGTAACTCA
1151 ATATATAAAG AGTTATATAA TACAACAGAG TTGATATATA TATAAATGTT
1201 GTATATAAAC ATAATATATA CGTTAATATA TATTAACAAA GAGTTGTATA
1251 ATACAACACA GAGTTAATAA TATATAAATA CAACACAAAG AGTTATATAT
1301 GTGTGTATTA TACATTTAAC TTAATATATA ATGAGTTAAA TGTATGTCTG
1351 TCCCATTCAA CTCTCCATTG AGGAAAGTAC CATTATCTTC CCCAAGTTCA
1401 GAAGAAGAAA ACAGAGAAAT ATATTGAAAT TCAGCAATTT GCTGGTGTGG
1451 TCAAGTCCAA CCCAGAACTT GCTTCTTTTA CATTGTAGTA CCCTCCAGGG
1501 TATGCAGAAA CAGATAGCTA GTGCATCTTT ATGACTAAAA AAGAAAATTT
1551 TTGTTGTTGA TTACCCAGTA ACAACAAGAC AGTATAAAAT CAGCATATTT
1601 TCTCAACAAT ATTTTCATTT TATAGTTGTT GAATAAAGTA TTGCTGACTT
1651 CATTTTAAAC TTTTCTACAT ACTTTGAAAA ATATGTTGCT TTCCTCCCAT
1701 TTTGTAAGTC TAGGTCTGCT ATTGATGAGC CATGCAGTGT TTTCTCCTGT
1751 TGCTTGATGT TTTTATTCTG AAATCATGGT TGGTTTTCAA ACACAAAAGT
1801 TTTCACTACA GTGATACAGA TGAGGTTTAT GTTTCCGCCA CAGTCTATAC
1851 TCAGGGTGCC TAGAGTATAG CATATTATTA GGGTACTATT TCTTTTCCTA
1901 TCCTAGATAT CCAACTAAGG CTTCGGGACA TGTTTTGAGC GAAGATGGGT
1951 GTTTCTGCCC GGATAGTATA AATCGAGGAT CCAGGTCTGG GCAGATTCAA
2001 CCATGGGAGC GAACACTTCA AGAAAACCAC CAGTGTTTGA TGAAAATGAA
2051 GATGGTAAGA AATATGGGAT AGTGGCATAT AAAAAATAGA ATTTTGCAAA
2101 ATTCAAGTAT ATGCTTCTAG TTTCATAAGT TAAGCATAAG CATGGTCTGT
2151 AGGGCCTTGA AGGAAAAAGG CAAAGCTGCA TGAGTGAGTC TGAGGACTTT
2201 GTAGGCTCAT AGCTAGGTTT TACCTTCCAC TTTCCATGGG ACCTTTGGCA
2251 GCTTTCCTAA TCTCCACTAT ACCAATGTCC TTTGTCCAAA GGGAGCTGCA
2301 GTTGGGCATG TGGTGGATAG TTAAATGATT TGTTTGTCCT CTGTGCTGTT
2351 CCTTGGCAGT TGAAGTTACC CCCATTGCTC ATTGTTACAG AAAATACATT
2401 ATCAACATGT ACATGAATGA TAACCAGTGC TCATAATATT ATAGAATGAA
2451 GCTGTGCCTT CTGAATTTCC AACTGCCAAG CTTTTGTGTA CTAGACAAAT
2501 CCCATAATGC TACGTCATAG AAAAAAGAAT CAGTTGTATT GGAGAAAAGG
2551 GAAACTTTCC AGGCCAGACT CAGCAAGACA AGAATAAAGG CATGAGTCCT
2601 CCTGATTCTC CCATCAGTGA GGCATGCTGG AACTGGGCAA TGCCTCCTCA
2651 TGTCCCTCTT CCTTCCTATA TGTTAAGTCT GAACAGCATT GGCGTATGCA
2701 GGTGGCAGCT GTTTATAGGT TGTCTGGGGG AAAAAAATGC CCCAAGCCCC
2751 AGGTAGTAAG TTGTCCAGAC CTCTGAGAGG GAGCTCTTCC GAGTAATTCC
2801 CAGAGAGCTC TGCTAATTGG AACAGGGAGG AAAAGAATGG ACTGAAATTC
2851 AGGAAATCTG ACACCAGTCC TACTACCAGT TACTTGCTAG GCCCAAGCAG
2901 CTTATTTACT GACTCTATCT TCAATTTTGT TATCAATAAA GTGAGGAGAT
2951 AGGTTCCTTC CCACTCAAGA AGTTTATCAT TTTGAGATCC TAAAGCAACT
3001 TTGTGAATTC TGAAGAAGCT TCTAAATCAT CAAGGAAAGT TTATTGGGTT
3051 AGAATGCAAG TTTGATTGCT GAAATGAAAA CTACAAATAA CAGTGGCTTA
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FIGURE 3A

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3101 AGCCAAATGG AAATGTTTAT CTTTCTCATG TGACAATCTA GGCATAAGTA
3151 ATCCAGGTGA TGTGTGGTTC CAGCAGCTTA GGGACTCTGA CGCCAACTAC
3201 TTGCCTTTTT CCCTCTCTTC CCATTTCTAG AGTGGTACCC TCAGAGTGGC
3251 TAACCAACAC AACAAATTCC AGCCAGTGAG AAAGGTGGAA AGTAGGAGAG
3301 GTTATGCCCA CTTATTTATA GGATTTGCTC TGGCTTGTCA CTTTCGTTCA
3351 CTTCCACTTA CCTAGATACA AGAAAGACTG GGAAATTCAG TTTGTTATCT
3401 TGGGTGGCCA TGAACCTTCT AAAAATAAGG AGTTCTGTTT TATTACAAAA
3451 GAAAAGAAGA ATTAGGAGTT TGTCATGATT GGGGACAACT ACGTCTGCTG
3501 TAGTTGGGGC AAACAATCTT AGTTTTGAAT CTTGGGATGG AAATACTTTT
3551 AAAAACAAAA TATGGGCCAG GCGCGGTGGC TCACGCCTGT AATCCCAGCA
3601 CTTTGGGAGG CCGAGGCGGG CGGATCACGA GGTCAGGAGA TCGAGACCAT
3651 CCTGGCTAAC ACGGTGAAAC CCCGTCTCTA CTAAAAAATA CAAAAAATTA
3701 GCCGGGCGTG GTGGTGGACG CCTGTAGTCC CAGCTACTCG GGAGGCTGAG
3751 GCAGGAGAAT GGCGGGAACC CGGGAGGTGG AGCTTGCAGT GAGCCGAGAT
3801 CCGGCCACTG CACTCCAGCC TGGGCGACAG AGCGAGACTC CATCTCAAAA
3851 CAAACAAACA AACAAACAAG CAAAAAAACC CAAAATATAT GGCTGATCAG
3901 GACGCCTTGT TTCAAGCTAT TCACTATCAG TTTGGAGGCC CATTCTTACT
3951 ATTTCTACAG AATAGTTCAT AGGAACTTTG AAATTATATA GCTGGAAAGG
4001 GGTCTTAAGA AAACTTTTTT TTCATGGCTA TTGTGATTGC CTTGCTTTAA
4051 CTTATCAAAT AGTAAAAGCA AAGATCTAGA GACTAGTGAT ATTACTTAAT
4101 TTTTCTGTCT CTAAAATGGA AAGACAAATA GGCTTGCTTT TCATTTAGTT
4151 GGTTTCCTCT GCTTCCTCTG GACTCAGAGC TAATGTTGTA CATGAGGCTG
4201 GTCGTCAGAG AATAGGGTGG AAAAGAGAGG CCAGCTGCAT ACTTTTAACT
4251 TGCTGGGCTA CATTTGAAGG TAGTAGAATA GCATTATGAT GAGAAAACAC
4301 AGAAATGCAT AACTCTTCCT TGATTCAGCC AGGCTTTGTT CTTGCGGGAT
4351 GCCCAAGAAA GCTACATAAC CAAAGAATTG TGACAATTGG GAAATAAGAT
4401 ACCCCTTTTT AGTTACTTTA AAGGACTCTA GAAAAACTAG GTTGAAGGAG
4451 AGTTAGGCTT AGGGACCAGA CAGGTCTTTC TTAACACCCT CTAGGTCACC
4501 ACCTTTTCTG TTGTCTGGCT TCTCAGCCCA ATGAGATGAA CCCACTGCAG
4551 CACCCATAAA GGAAAGATCT GAGCATAGCA ACAAGTCTGT GCCTCCCAAA
4601 GGTGCTAGGC TCTCTGTCTG TTTATGCAGA CAGTTGCAAG GCAAAGGAAG
4651 TAGGAGGGCA AGTCCACCTA CTATAAACCT GTCACTCTCT AGACATGAAG
4701 AATAGAGGAG GAAACAAGTT GGTCCTTGCT CTGTCATTGT GAACCCCATG
4751 TTCTGATGAT GGAAGGCTGA CAATAAAAAG GTAAATAATA CATAAACCAG
4801 ATAATTTCAC AGTGCCTTAA AGTGCCACCA AGGAAATGAC TCCTAGTGAT
4851 CTTACAGACA GTGACAGTGA TGGTGAGGAG GCCACTTTAG ATAGGGTGGC
4901 TGCGGTTGTC TTTCTAAGGA GGTGACATTT GGGCTGAAGC CTGAAAGATG
4951 AGAAGAAGCC ATCTATGAAA TGACATGAAA AGAATAGTTC AAGAACAGGA
5001 AAAACAAGTC CAAAATCCAA ATAATGACAA AATCAGGATT GAATAGTTGC
5051 CTATATCTTA ACGTTCTCTC ATGAGCACTA GTTTGCCAAA GAGACTGCAT
5101 TTATTGCCAT GTTAACTTAT TTCTTCAAAA GATGATTGAT TTGAGGAGAA
5151 AAAGTATGCC ATTCTAGGGA ATTTACTTTG CTTTAAAATT CAGTACATTT
5201 TGTAAAGTTC ATTTGACTCT TCACATAAAT CTGGATTGAG CACAAGGTAA
5251 AATTGTATCT GATTGCTGTG AAGCTCCTGA CCAAGAAAAA GCAACCAAAA
5301 AGCACTGATT AACCAAACAA CATTAATGCT TATGTCATTT TTGATATCCA
5351 TATTTTTATA TACATAATCA TAATGTATAA TCAAACTGGG CCAGTATCAA
5401 GGGCACTAAA ATGAGCCAAC TTAATTATTT AAAAAATATT GCTGAAAAGA
5451 ATCCCAATAT GTGATTTTTA AAAAGTTTTT TAAAATTTTT AAAAAGATTT
5501 TTTAAAAGAT TTTTAAAAAT ATTTTCTTCA AACTGTTTAA TATTTCCAAT
5551 ATATAGATAT GAGAAAAACA TTTAACCAAT AATTTTCCCA AGTAATGTTT
5601 CAAGAATTCT CTCTTATGGA AAAAGTGTTT TTGTTCACTT TGAAGGTAAT
5651 TAAGGAGCAA GATAAGAGGT TATTGGATGT CCCTTGAGAT AAGCTATTCT
5701 TGCCAGAATT CATCCTGACA CTTGTATTTC ATGTTGTTCC ATCTGATATC
5751 TGATCTTGAA CACATAATTT TATTAGTTAC TTATGTTGAT CTTTATTCAG
5801 CAAAAACAAA GTAGGAGATT TTCAGGCTAG GCATGGTTGC TTACGCCTGT
5851 AATCCCAGCA CTTCAGGAGG CCGAGGCGGG CAGATCACGA GGTCAAGAGA
5901 TCGAAACCAT CCTGGCCAAC ATGGTGAAAC CCCATCTCTA CTAAAAAATA
5951 CAAAAAAAAT TAGCTGGGCA TGCCAGTGTG CGCCTGTAGT CCCAGCTATT
6001 CAGGAGGCTG AGGCAGGAGA ATCTCTTGAA CCTGGGAGGT GAAGTTTGCA
6051 GTGAGCTGAG ATTGCTCCAC TGCACTCCAG CCTGGCAACA GAGCAAGACT
6101 CTGTCCAAAA AAAAACGGCT TGCTTATTTG ATTATATAAG ATATCTTTCA
6151 TAAATTAGAT CTCAAATTAT ACTATTGTTT TGCAGTTTTA GCTTTTATGT
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FIGURE 3B

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6201  TTTAGGGCAA  ATCTTAAGTC  CTAATTACTT  TTTTTTTATT  ATTGTGGTAA
6251  AATGTATATA  ACAAAATGTA  CCATTTAATC  ATTTTAGAAT  ATACGGTTTA
6301  TGACATTAAG  CACATTCACG  TTATCATGCA  ACCATCACCA  CTACCCATCC
6351  TCAGAACATT  TCTCTTCTCG  AATTGAAACT  TGGTACCTCT  GAAACAATAA
6401  CATCCACATT  CCATCCCCTC  CCCAGTCCCT  GTTAAACAAC  CATTTGACTT
6451  TATGTCTCTA  TGAATTTAAC  TACTCTATGT  ACCTCATATA  AATGGAACAT
6501  ATAAGATTTG  TTCTTTTGCA  TCTGGTTTAT  TTCATTTAGC  ATATATTTTT
6551  AAGGTTCATC  CATGTTGCAG  CATGTGTCAA  GATTCTCTTT  CTTTTTAAGT
6601  CTGAGTCGTA  TTCCATTGTA  TGGATATACC  ACATTTTGTT  TATCTTTTCA
6651  TTAGTTGACA  TTGATTGTCC  TCACCTTTTG  ATTTTTGTGA  ATAAGGCTGC
6701  TATAAACATT  GGTGTGCAAA  TATCTGTTCA  AGTCCCTGTT  TTCAATTCTT
6751  CAGGGTATAT  ACCTAGAAGT  GGAAGCACTG  GATCATATAA  TTCCTTGTTT
6801  GACTCTCTGA  GGAACCATCA  TACTGTCTTC  TACCTAATTA  TGCTTTGTGT
6851  TTTAGTAATG  GGACACAGCC  TGGCATGATG  GGCTAGAGTA  TTGGAAAGGC
6901  ATGCACAGGT  TCAAGTCTCA  GCTGTGCCAC  GTGCCAGTAA  TCTACATGTT
6951  TCTATGAGAA  GAGTCAAAGA  GGATATAGCC  TGGTCAACCA  TTATCAGACA
7001  CTGGAGTCAG  TTTGACTAAT  TATATGGTGT  TCTAAGGAAA  CTTGAGGTAC
7051  CACAAGAAAA  GTCTCCAAAT  CTAAATAATT  ACTAATGAAT  TAATTGAGGG
7101  GGAAACTTAT  TTAACCTTTG  TAAGCCTCAG  TTTCTTTGTA  TGTAAAATGC
7151  AGGTAATAAT  TGGGCATACT  TCATTAGGTC  TTTGTGAGGA  TTGAATAAAT
7201  AATGCAAGTA  AAACACTTAG  CAAAGTATTT  CCCATAAAGT  AACCACTCAA
7251  TTAATGCTAA  TTAAGTGTTA  TTTACTAACA  TCAGAGTTTC  CTAGTGTGAA
7301  CTCTTTGAAG  TACTTTAAGT  TCTGAGAAAA  ACAAAATTAA  TTAAATGCAA
7351  CTCTGTCGAT  TCCACAGTTA  ATTAGACCTA  TTCATGTTTC  TATTGACTGG
7401  ATTAACAGAA  CGGCAGATTT  TATGGATTCT  GTTAAAACCT  ATATAAAAAC
7451  ACTTTAAAAG  AAGCCAAGTT  ATTGACTGCA  CAAAAACATA  ATCTCATCTG
7501  ATATCTTTTT  TATCCCCCTG  AGGTTATTGT  GTTTTTGTTT  AAGGCAAAAT
7551  CAAGAACTAA  TTGGGATGAA  AATAACTAAA  GTTTACTTTG  TCTGATTTAA
7601  GTCCCAAACT  GACTAATAAG  TAATCCCATT  TGATCAACAG  ATTCAGTGAA
7651  AACTGTCCCC  CATTCTCAAC  TACCATATGG  ATATTCTGAG  AAATAATTAA
7701  TGATGCAGAA  AAACATTTTT  TGTTTTCTGA  AATAAAAGAA  TAGACGTGCA
7751  AGTGACACTT  CTTTTTAATG  CTTACAACCT  TTTTTTAAAA  ATCTACTTTA
7801  TTTTCTCTAT  CTGAATGCAC  TAGATTTTGT  TTGTTTGTTT  TTGTGGTTGG
7851  TTGGTATGGT  TTTGCTTATT  GAGGTTTTCA  GGCTGATTTA  GAAAAAAGAA
7901  ATTTTTACAG  GAGAGAGTGG  ACTTGTTTAC  AATTCAGAGT  TGAGGCAACA
7951  AAAAAAAATC  TTGCAGTCAT  TATGAGTAAT  ATGTGTATCC  AAGTTTATAC
8001  AAAGAATGTA  AAGGTGATAA  AGTTGGCTTA  GTTAAATCAA  GAGACAGCCT
8051  TCTTCTAGAA  TATTATAGCT  AAGAAAATTT  GGACTTAAGT  TTAAAAAGCT
8101  GCTCTAAAGA  GTTCATCAAT  GCCCTGAGTT  TGCAGAGAGT  TCAATTATTG
8151  CATTATTCTT  TGGACTTGCT  GAAAACTCAG  TGTTCTACTT  TTATTTGGCA
8201  ACACCATCTC  CTAGGATATG  TGGCTGTTTC  CAGTTTTCCA  GCATCTTCAG
8251  TGACAGAGGC  AATGGGATCC  TTTAAAATGT  TGGGCCAAGA  AAATTGGCCA
8301  CAGATTTGCA  ATCCAAAAGA  AATAGGAGGT  TGCTAAATTG  ATTCCAGCTA
8351  TGAAGGACAT  CGAAAATTTC  TTTTGTTATT  TGACTGTCTA  TCATGGTCTA
8401  TTTGCACTCA  ATTTAATAGG  CAAATGAATT  TCCGACTTTC  CCTTAGCAGC
8451  CTTGAGTAAT  GCTGTCTCGT  ATTTATTATT  TTGCATTAGA  ATGGTTGGAA
8501  AAGTTAAAGG  AAAATTTCCC  TAGCAAGAAT  TGGCTTCTTA  AAAAAAATAAG
8551  TCATCTTGGA  CAACCTAACA  TTTAGTAAAG  GCATTTGTCA  TAAATAACCT
8601  CAAGTCCAAT  TTATGGCAAG  GGTTTTAATT  TGTAAGGGCT  TTATTTCTCC
8651  ATACAAAGGG  ATTGGAGAAA  CAAACTAGAA  AGCCAGAAAA  CAGACCACAA
8701  ACACTGAGCT  AGTGGTTCCA  ACTGGAGTGT  TCCCTGAGCA  GTGACTTATG
8751  AATACTTGTT  TAGAAGAATC  AACTCAAACA  AATTTAGGAA  AGTCACATCC
8801  TGCCTTTAGA  GCTTCCAGTG  TTTGTTAGCA  TATTAAAGTC  TCTGAAATGA
8851  CCTACAATAT  TGAAATCTCA  GTCTTCTGCT  ATTTTTAATA  TTTATTTCAA
8901  AATGAAATAA  TTTTTGTGAA  AAACATTTTA  ATGTCTGTGG  CTCATAATAT
8951  TCTGTGGATC  TCAGTTTGGG  AAATGAAAGA  TTATAATCGT  ATCTACTCTT
9001  TATCTGTTGG  AAACATCTTT  CCATTTATTT  TTCCTGCTGG  TTTAATGGCA
9051  ACAAATTTTT  ACATGTGAAA  TATTTGTAAT  GTGATTTATA  TGAAAAAATG
9101  TAATTTTCTT  ATTACACGAT  CAAAAGTGGT  TATGCTCCTC  TGTAAGTTTT
9151  TCCTTACAAG  TTTTTATGTT  GCATAATTTA  TATCTATTTG  GTTTAATGAG
9201  TACAACACAA  GATAGCTCAG  TTTAATTCTG  GGATGTTGGA  TGTTTCTAGT
9251  TAAAGTACAA  GTTGGATTTG  ATGAAAATTC  ATTGCTTCTT  TATGATTTTT
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FIGURE 3C

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 9301 TAAAACTCAA GAACATGTTA GTTAAAGAGT GTCTTCTGAA CAAATTCTTG
 9351 TGAAGTAGTT GCTGATTATT AAGTAACACT CATGCTACCG TAACTTTTTA
 9401 TACTATCCAA AGCTATAGAC ATTTTTAATT TTCAACTTGC AACTACCTAG
 9451 GTTGAAAAAT TAAATCTGCA AGCCAGTTTC ATTATTCAGA CAATTTGGTT
 9501 ATCACTTCAA GCCTACTATC TTCAAAGAAA ATGGGAGTGC AGGCCTTCAT
 9551 GGGAGCTGAC TTCTGCTGTA TGGCCTTGCA AATGTCAACT CGATTAGAGT
 9601 GACCAGTGTT AGCCCTCAAT TCACAAACTC AGGTCCCATG AAATATACAC
 9651 GGATTTCTAC TATGCATTAC TATGTGACCA TTCATGGAAG TTTCGTTTGG
 9701 AAACACAGAC ATTAAAAAGC CAGTCATGGA ATAACATTCT TGTTAAAACA
 9751 GGACATTGGC AAAAAGGACT AGAAAACTTC TGGCTATAGA TTTTGAATCC
 9801 AATAGCCTTG CATAGGCTTT TCTGTTTCCT CCTAAACTAT GTCTTCTGTC
 9851 CTTTCTGGAG GCATATTTAT AGTAAAATAA ACAAAATTAA CCTTGTTTTA
 9901 CACTTGAGTA ACCTATACCT TTGGTTATTT ACGAGAATTA CTTAAAGCAG
 9951 AGTTGGCAAC TTTTTCTGTG ATGGGCCTGA TACTAAATAT TTTACACTTT
10001 CCAAGTAATA CAGTCTCTGT CACAACTACT CAACTCTGCC ACTGTAGCAT
10051 AAAAGCACAC TTAGACAATG CAGAAACAAA TGAACATGGC TTTGTTCCAA
10101 TAAAACTTTA TTTATGGACA CTGAAATGTG AATTTCAAAA ATATTTTTTG
10151 CATAAGATCA AATATTATTC TTTTGATTTT TTTCCAATCA ATAAAAAGTG
10201 TAAAAATTGG CCGGGCATGG TGGCTCATGC CTGTAATCCC AGCACTTTGG
10251 GAGGCCGAGG TGGGCAGATC ACCTGAGGTC ACGAGTTCGA GACCAGCCTG
10301 ACCAACATGG AGAAACCCTG TCTATATTAA AAATACAAAA TTAGCTGGGT
10351 GTGGTGGGGC CTACCTGTAA TCCCAGCTAC TCGGGAGGCT GAGGCAGGAG
10401 AATCGCTTGA ATGCAGGAGG CAGAGGTTGC GGTGAGCCCA GATTGCACCA
10451 TTGCACTCCA GCCGGGGCAA CAAGAGCAAA ACTCCGTCTC AAAAAAAAAA
10501 AAAAAAAAAA GTGTAAAAAC CATTCTTAGT TCATGAGCTA TACAAAAATA
10551 GATAGTGAGT TAGATTTGGC CCATGGGGCT TATTTTGCTG ACTCCTGCTC
10601 TAAGCATCTT GCAGACATTT CTTCATATGC CCTAGGAGAT TTCTGATATC
10651 CCCTCATAAT ACCCTGGCCT TACACCAAGA CTACAATCTG TTCTTTGCAG
10701 ATGCTTAATA AATTCATTCT TCCCTGTCAT TCAGTTGATC TGTGTGAGCC
10751 AGTGGAAATA CTTGGGCCAA TAAATCTAGT GTGTTTGAGG GTAAAATATG
10801 CTATTTTTGT AAGATATATT ATTTAATGGC CACACAACCT AAATTCAATT
10851 AAATGGTTAC AACCTGTAAC GCATTTAAAA TATGACTAGG CAGAATTTGC
10901 TTCCTACTAA AGACATTTAT TCGATTGAGG AGCATCCAAC AGTTGATGTT
10951 GATCCCCCCA TCCTGCCCCA CTGTTCTACT TTGCAATTTG TTTGAAAGAA
11001 ATTGTCAATA TATTTCTGAC TTCTGAGCAA ATCCATGAAT CGGGATCCAG
11051 CAACAGGAAA AGAAGCTGTT GCTGCCCATT GCTTGGTTTT GGCACCAGGA
11101 ATGGATAAAT CCCAGACTTC CTGGGGCACG TGTTTTATAA AAGGGAAGTG
11151 CTGACAGTGC AAACAGCTGC CATCAATTGG CCTTGGAGAC TACTTCCCTG
11201 GAGAAGCTCC AATTATATTC TTAAAGGACC CACCAAGCTC TTCAAGTGTT
11251 AGTGGCAACC ATTTGCTGCC AACCATTTGA AATGATGAAG TAATTTTTTT
11301 TTATTAGTGG ATCCTAAGTG ATAGGCTCTA GAACTGATCT TCAACCTTAA
11351 CTAATATCAT GGCATCAGAG GGCTACAGAT TAAATCAGTG GTTCCCAGTC
11401 ACTCTCTGTG GACAAGTAGC AACTACGACA AAGCTTTTCT TAGTCTATGG
11451 TGGAAGAGAA AAATTAGGAC AATGTAATAA GCATCCCATA AACTTATTAA
11501 ACCTATTAAA ATTTAATTTT AAGATTATGT CATTTTTTGT ATGTGTGTAT
11551 GCTTAGTATT TATGGATTGT GGAAATAGAA TTTTTTTTTT ATAGTGAGAA
11601 CCTAGGTAAG TGACTTACCT CTCTGATCCC CCATTTTCTC ATATGTAGAA
11651 GGGGGCTAAT AATAGTATCT GTCTCATAGT TTTTGTGAGA ATAAAAAAAT
11701 TGTCCAGGTA AAATGCTTAG CTGGTGACTG GCACACAGTA ATTGCTCAAT
11751 AAATGTTAGC TATTATTGCT ATCATTATAT AATCATCATG GTTTCCAATG
11801 CCTTTACTTG GCAAATAAAA GAACAAAAGT CACCCGATAT TGATCTCCCT
11851 TTTCTTCCCT AGTTTTCTGG GGGGTGGGAG GCAGAGACCG AATTTTCTGA
11901 TCTGTGAAAT CTGAATTTAT CATTGTAATT TTCCATAAGT GCTATGTAGA
11951 GAACTCATTT AAGTTGCTGG GATGAAAAAA AATCAAAAGT GGCCTATTGT
12001 GCTGGGTGCA GTGGTTCACG CCTGCAATCC CAGCACTTTG GGAGGCTGAG
12051 GGGGGTGGAT CGCCTGAGGT CAGGAGTTCA AGACCAGCCT GGCCAACATG
12101 GTGAAACCTT GACTCTACTA AAAATACAAA AATTAGCCTG GCATGATGGT
12151 GGGCACCTGT AATCCCAGCT ACTCAGGAGG CTGAGGCAGG AGAATCCCTT
12201 GAACCCAGGA GGTGGAGGAT TCAGTGAGCC GAGATCTACT GCACTCCAGC
12251 CTGGGCAACA GAGTAAGCCT CTGTCTCGAA AAAAAAAAAA AAAAAAAAAA
12301 AAAAAGTGGC CTCATCTTCA TTTCAGTGAA AGATGATAGT ATCTGGACTC
12351 ACAGTGTGGC AGTGCAGACG GAAAGCTGAG AGTTTATTCA ACATTTATTT
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FIGURE 3D

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12401 TCAATATAAA ATAATTAGGT GTTACTGATG GCTTGAATGT GGGGTAAGAT
12451 GGAAAGAACA AAATCAAGGA TAAATCCTAG GTTTTTGCTT GAGTAGTTAT
12501 GTGGATGACT GTGACATTTT ACTAAGATGG AGATGCGTGG GAACGGAGGG
12551 GTTTGGGACC CTGCTCACAT ACAGTCTAGA GTTCACTTTT GGAGGCATAC
12601 AGTGATTATG GGACAGCTAA ATGATGGTGC CAAGTAGGAG CTGGAGTAGA
12651 GTATCCAGCA ATGAGTGGAA ACATCTGGGA TGGAGACAGA AAGACACGGG
12701 TATTAATTCT ACGGGGATGG CTAAGTCTGC TCTGAGAGAC AGTGTGGAGA
12751 CCAAGGAGAA GAGGAATCCT AATATTTAGA AACAAGGCAG TGGATAGCAA
12801 TCTAGCTATG GAAAGTGGAA GGAAAGAGAT AGTTGATCAT CCAGTTCAAC
12851 ACTACTCTTG TTGTAGTTCA CTTATGTTGA ATGCTTCTGT GTGACTAAGT
12901 CGGTGAGAAA AATCTATGGG AGTAGGCAAC ATGGAGGATG TTGGTATTCA
12951 CAAAAGCAGT TTAGTGGAGT GTGGAGGCCT GAGCCAGACT AGAATGAGTT
13001 AGGAGTAGAT GGAAGATAAG AATGCAGATA TGGGCCCAGC GCGGTGGCTC
13051 ACGCCTGTAA TCCCAGCACT TTGGGAGGCC AAGGTGAGCA GATCACAAGG
13101 TCAGGAGATC GAGACCATCC TGGCTAACAC CGTGAAACCC CATCTCTACT
13151 AAAAATACAA AAAATTAGCC GGGCCTGGTG GCGGGTGCCT GTAGTCCCAG
13201 CTACTCGGGA GGCTGAGGCA GGAGAATGGC GTGAACCCGG GAGGTGGAGC
13251 TGGCAGTGAG CCGAGATGGT GCCACTGCAC TCCAGCCTGG GCAACAGAGC
13301 AAGACTCCAT CTCAAAAAAA AAAAAAAAAA AAGAATGCAG ATATGGCAAG
13351 TATAGACAAG CTTCAAGAAG TTTGGTCTAA AAGGAAGCGG AGAAATAAAC
13401 AAAGAGATGA TGCCTAATAT AATTCAGCTA AATGTAATAT AATGGATTTT
13451 TTTAAGATGA GGTACTAGAG CATGTAATAT AAATCTATTA AATTGGGTGG
13501 CCAGGAACCA GGACTGGCTC ATCAGCATGG ACCAGGCTAG ACGCACAGGG
13551 CCTTATATCC AGAAGGACAT CACCTTTGGG TTTTAATGCT CTGCACTTGC
13601 TGTCTCCAAA TTCTAACTGT CTCTTAGGCT CTCATCAACA CCCACCTCCA
13651 TATCCAGATA TTGAGTACCT CAGGGAGTTC AATTTGGAAG CAAATGATGT
13701 GAAAATGTAC TTTACTATCC AGTAACATTC TTGTTAGGGA GTGTTGGCAG
13751 AGATTGTCGA ACAACCATAA TGCATTTTAT CATTCGATCA GTCTACAATT
13801 TAAACATAGC AGGACTGGAC AGAGGCACAG GAAGATTAAG CCACTGACCT
13851 TAAGTCAGAC AGTCACATGG GTAGATCCGG AATCTTGATC TAAAATGAAT
13901 ACCATTTTTT CAGTTATAGC TATCTTCCCA GGATGGCCAA CCAGAATGCA
13951 TATATAAAAT TTCAAAAACA AACATTGGGA ATTGCTCTTC AGCAAGAATA
14001 CATCAAACAC CCATTATGTG CCTAACTCTA AATCTTACTT TCAGAGAGCT
14051 AAAAACAATT TCATTTCACA GTGACATTCA TCTTCGCTTC TGCCGTAACT
14101 CACATGCATA TGCCTTAGAC CACATTATTA ATGAAGTATT GGGGGGGTTCC
14151 ATCTAGAGCA CCTTTTCTTC CCTGGAGTTA ATCATCCAGT TCAGCACCAC
14201 TCTTGAGCTT TGCTTAGCTT CTTCTACCCA TTTGGATTTT AAGGACAACA
14251 ATTCCAATGG CCTTTATCCA TGTATTTAAC AATTCATTAT GAGCCAGGTG
14301 AAGTGGATCA CACCTCTAAT CCCAACACTT TGGGAGGCTG AGGCAGGTGG
14351 ATCGCTGGAG CCCAGGAGTT CACAACCAGC CTGGGCAACA TGGTGAGACT
14401 CCATCTCTAC CATTTTTTTT TTAATTAGTT GGGTATGGTG GCAGGAGATC
14451 AAGGCTACGG TGAGCTGTAA TTGCACCACT GCACACTAGC CTGGGCAACA
14501 GAGCAAGACC CTGTCTCACC AAAAACAAAA ACAATTTATT TCATCATCAT
14551 TGTCATCATC ATTGTCACTG CTCACTCTTC AACATTTTTT AGGTCAACTT
14601 AATTAATATG ATACCTTGTG GGATAATTTT TATTTATTTT TATAAAATAT
14651 TGAAGTTTTT GCCACTTTGA TAACTTCTTC ATTTTCTGTC CAGAGTATAA
14701 CATACCAGGG AAAAGGCTCT AAAATAAGGC TTGAGGTATT AAAAAGATCT
14751 TCTGTTTAAG TCTTATGTTC CTAATCAATA ACTAGAATTG GCCTGATTGC
14801 TTTCCTCAGT GGGTTTTCTG GTAGTCCTGA TATGATATCG AGGCTGTCAT
14851 ATAGTCCTGA AATATCCTAT CATTAACATT TGTGGTGGTA TCTGATATAA
14901 AGGTAGATGA ACTTCATTGC AGCTATTCTT AGGAAATGCG TATTTAAATG
14951 CATAGTTAAA AGCAAGATTT ACAATTATAG AAGGAATGCA AATGAGTTGT
15001 AGAAAGCTCA TAAAATAAAA ATCAAGAAGA AAGAATTACC CATCATGCCT
15051 CAGCCCAGTG ATAACCACTG CTAATATTTT TGGCTGTTTT CATTTGCAAC
15101 CCCATCTCCA TTCTAGCAGC CCTCATCCCT CCTACCCACT ATGTTTTTCA
15151 CTATATTTCT TGTTTAAATT TACTTAATTA TTTGTTAATT ATGTTTTTCC
15201 TCTCACTAGA AAGTGAACTC CATGAGGGCC AGGGATTTTT GCTATTTTGT
15251 TCACTTTTGT ATCCTTAGCA CCTACTTTGT TGATTAAGTG AATGCATTAA
15301 TGATCTATTT TTAATCTGTG TATGTGTATA AAAGACACTT GATATATCTG
15351 GGATGATATT CAATATACTT TTGTATCCTC ATTTTCACCA TAGGTAGTTT
15401 ATGTCAATTC CTTGAAATTT GTTGATTTTC TTGAATAATT TAGCAGTTGT
15451 ACAATTCTAA AACATAAATA TAATTTGCTT AAATATACAT ACCATTTTAA
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FIGURE 3E

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15501 ACATATTTAA ATGTGAAAAT ACAGTTGAGT TCTCTTAGAT TGCAATTTTG
15551 TAACTTTTGA TAATCCTTTG ATCCTGAAAA AAATTTTTTG GCATGAGGGA
15601 AGAGATGAAT ATTTCTTTTG GAGTATTTAA ATCATCTCTG CAATAATCCT
15651 TTGATCCTGA AAAAAAATTT GTGGCATGAG GGAAGAGAAG AATATTTCTT
15701 TTGGAGTGTT TAAATCATCT CTACAATTAA TAATATCTAA AGCAGTTTGG
15751 TTGGTTTATT TAGGTAGGAT TAATTTTCAG TATGAATATT ATTTAAAAAA
15801 CAAATATAGT CAGTTGAATT GCTGTGGAGG TTTCTGTACG ATTTACTCAA
15851 AGCTGGCTCT TTTTCTGTAC GCACTACCAC GCCCGGCTAA TTTTTGCATT
15901 TTTTTGGTAG AGATGGGGGT TTCACCATGT TGGCCAGGCT GGTCTTGAAC
15951 TCCTGATCTC AAGTGATCCA CCCACCTCAG CCTCTCAAGG TGCTGGGATT
16001 ACAGGCATAA GCCACCATGC CCAGCCTGCA TTTATCCTTA CATGATGGTG
16051 AAAAATAATG TTTGTACTTC CTTCAGAATA ATTTCAAGAA GGATCCCTGG
16101 AGTCAGCTAA TGATTAGAGT CAGGACTGTG CCTTAGTTGA TGGCCCATAT
16151 AGCACTACTG AACATGCCAG AGCTTTTGCT TATCCATACT GGAGGAGGGA
16201 GTGCTTAGAA GGCAAACGTA TATCATTTTA TTTTCATTCA AAATGTACTG
16251 ATAGCAAAGA ATTTCAATGG CTGGCAGATT CAGTTAAGGA CAAAAATAAT
16301 TCACAGCAGA AACTTTTTCT TGGTCTCCCT CCTCCAAGTG CTAAGCATGG
16351 CACAAGTAGA TATCATGGAA TTCTAGAACC CTCTCTTCAT AGATCTTAAA
16401 AACTACTCTC TTTCCCTGCT TGAGTACTTT CTCAAATCTG TGTCTGTGTG
16451 CAAATTTTCC TTCTAAGGAC ACCAGCCATA CCGGATTCAG GGCCCACTCT
16501 ACTCCATTTT GATACTGTAC CATCTTAACC GAACATGTTA TATCTGCAAC
16551 AACCCCATTC TCAAATAAAT TTCACAGTCT GACATACTAG GGGTTAGGAC
16601 TTCAACCTAT CTTTTTGGGA GACACCTTTG GTTTGACTGC TTCTTCAACT
16651 CTTACCAGCT CTATGAGCTT GAGCAGGTTA CATACTCTTT TCAAGTCTTA
16701 GTGCTTCACT TGTATTTTGG GGCTAATAAG GATTATACGA AATAATGCAG
16751 GTTAAATGCC TAGCACTTTG CTTTACATAC TAAGGGTTCC CAAGTGCTTT
16801 ATTATTAGGT TTCTGAATGT TATATATAAA GTTTCAGTGC TGCAAAAGGA
16851 ATAGCACTCG AATATAACAT TTTCTTTTTA ATTCTCAGCA AGGCAACGTA
16901 CTTCTATATA GAAGGGTGCA CCCTTACAGA TAGAATAATG GTGGGCGCAC
16951 ACTTGGACAA GGGAGGAGAA GGGGTTCTTA TCCCCCACGC ACGTGGCCCC
17001 TGCTCCTGTG TCGTTCCCCT ATTGGCTAGG GTTAGACCAC ACAGGCTAAC
17051 CTAATTCTGA TTGGCTAATT TAAAGAGAAT GACGGGGTGA GGGCTTTGGC
17101 AGAGTCAGGG CAGAGCAGAT AGCAGGTAAT CGGACTGAGT TAGGGTGGAG
17151 CAGGTGATCT GAATGAGTCA GGGTGGAGCA ATCAAAAAGG TTGCTTTATG
17201 AGGAAGTTAC GTTTAAAAGT AGAAGGCAGG CTGGGCGCGG TGGCTCACGC
17251 CTGTAATCCC AGCACTTTGG GAGGCAGAGG TGGGCGGATC ACGAGGTCAG
17301 GAGATGCAGA CCATCCTGGC TAACACGGTG AAACCCCGTC TCTACTAAAA
17351 ATACAAAAAA ATTAGCTGGG CGTGGTGGCA GGCACCTGTA GTCCCAGCTA
17401 CTCAGGAGGC TGAGGCGGGA GAATGGCATG AACCCAGGAG GCGGAGCTTG
17451 CAGTGAGGCG AGATCCTGCC ATTGCACGCC AGCCTGGGCG ACAGAGACTC
17501 CACCTCAAAA ACAAAACAAA AAAGTAGAAG GCAAAGAATT GAACATACTG
17551 ACATATTAAG TCTTTGAAAA GAAATTTAGA ACTCATATCT AACAATCCCT
17601 CCCCTTGTAT TTCCTTACAG CTTTCTTTTC AAACTTTTTT TTAATATGCC
17651 TTGGCTTAGT AGTTTTGCTT CATTTTCCAA AAGAAGAAGC TTCTCTGGAT
17701 AAGGTGGAGG TTAGTTAAGG GAGGTTTCAG TAAGTGACAT TTTTATGAGC
17751 CTCTGCATCT ACTTACGGAT GCACAGTATG ACACAGCACC CGACAAGAAT
17801 AAGTCCACCT ATTACGGCTG CGAGGGAAGT AAGAATTGAG GCTATTATTC
17851 CTTCTCATTT ACCAAACTAC TTTTCTAGCC ATCTTATAAA GGGGTCATTT
17901 ACCCCTGAGT TGCTGGCTAA CTTATTGGAT AGAGCAGTCA GACCATGCAG
17951 TGCCTTTCTA ATACTTCCAT TAGGGGCAGT GTTGTTTGGG ATGAAGGTGC
18001 AACATTGAGT TTTAATTATG ATGCAAACTA CCCCTCTTTC TGCTACTATC
18051 ATGTCTAAGG CTATTTTATT TTGCCAAGCC ATCTGGCTAG TAGCCCCTAA
18101 TTGCTCAGCT ATTCCATTAA CAGCATCTCT AGTGTAGTTA ATAAATCACT
18151 GTTGGTTGTA GTAGCTGTAG TTTATCCAAT CTACATTTTT ATTAATTGTC
18201 ACTCACCAAA ATATTGACTT AAATCCTGCG GCTATTTGAT TTTGGGCTTT
18251 AAATTGATCT GGTATTCCTC ATGGGACCCT AATTGTGTCT AAATAGACGT
18301 GAGAGTTGAA AGACCCATAA GGGGCTTCTC TCGCTTTACG ATGTCTTATT
18351 TTTCCTTCCT CTGGTTGATG AAATGCCAGG GTGAAAGGGA TAGCCAATTG
18401 GACTAAAGCA CAAGTGCCAC TCCAGTTATT TGGCAGAGTG TCCAGTAAAG
18451 GTCCACCACA ATACCACCAC ACATCCACAC ATCCGCTCGG GGATGAATAA
18501 GGGCTGACTG ATTGATAAGC TCTTGAAAAT TCTTAAGCTC ACTGCATCCC
18551 TTCAGGTCTC CAAGGAACGC TAAGTTTCCT CCCTGTCATG AGAGACACTA
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FIGURE 3F

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18601 AGTGAACTAG TTTTGGGAGA CAGAAGCTGG ATGGCCCTTG GGGGCTGACC
18651 TGCAGGGTAC CAGACTTCGG GATATAGCAG AGAGAGAGCT TGGAACGACT
18701 TATTACTCCA GGCTGTAGAA TCCCTGGAAA AGAGCTACCA TGCAGCCCAT
18751 GCCTGGTTGA CTGGAGGACC ACCCTAGTGG AAAGGGGACA ATCTGGAATA
18801 CTTGATCCAT TCTAACCAGG CATTTGCATC TTGGTATCCT GTCTTAGTTG
18851 CCAAAGTTTG CTTTAAGTCT TTGTTTTTTT GTTGTTTTGT TTTGTTTTTT
18901 GAGACGGAGT TTCGCTACTT GTTGCCCAGG CTGGAGTGCA ATGGCGCAAT
18951 CTTGGCTCAC TGCAACCTCT GCTTCCCAGG TTCAAGCAAT TCTCCTGTCT
19001 CAGCCTCCCG AGTAGCTGGG ATTACAGGCA TGCACCACCA TGCCTGGCTA
19051 AGTTTGTATT TTTAGTAGAG ACGGTGGTTT CTCCATGTTG GTCAGGCTGG
19101 TCTTGAACTC CCAACCTCAG GTGATCCCCC TGCCTCGGCC TCCCAAAGTG
19151 CTGGGATTAC AGGCGTGAGC CACCGAGCCT GACCTGTTTT AAGTCTTTAG
19201 TTTTTACAAT AGCTATCTTG GTCTTGTTGT TAGATGGAGG AGGAGCAACT
19251 GTTCCGTTGT GAGAGGTTTT GGAAGAAGGC TTACAGGAAG GTGCAGGCGG
19301 TGGGGATCAA AGAAATGCAT TTTAAATAAT CTAATAGGGT TTGTCCCTGA
19351 AACCTCAGCC CCTATAGCAT AAAACTGACT TAAAGAAGGG AACTGGCTTA
19401 GAAAAGGGGA AGAAATTTGA GAGTTTGAGA TAATAACCTG TAGAGAATTA
19451 TAGATAATAA CCTGTATAGG TTTAGCTGAC AGCTGGGGGG AGGGCTGTCT
19501 CTTTAGTAAA ATGAGTGTAT GGTTTTAGTA AATTACAAAA ACTGGTTGGG
19551 GCAATCCCTT CTTGCTATTT AGTGGTCCAC AGAACATTGG ACCAACTACA
19601 GCATAAAAGC TCTACGTCGG GGGCGGGGCG GGGGGTAGGA CTCTGGGTTG
19651 ACATTGGGGT CTTTATTGAA ATTTCCCCGG ATTAAATGGT CCCAATTCAC
19701 TAATGCCCAG TCTGATGACA GTCAGGAGGC ACAGAGGTAT TTTTTCTGAA
19751 ATAGAGAGGT GTCTTTGACT TGGCAAATCC CCACAGGGTA TAACAAGGCA
19801 AGCATTAAGT GCAATAGTTT GAGGCAAAAT TGACTTGGTT ATGTTAATAA
19851 CTAGATGGTC AGCAATAGAG CCAGTAAAGA AGAAAGAGTA ATAGAATAGA
19901 TAAAAGAGAG TTAAATTTTT CTTAGCTTTA GTTTGGCAGG GCTTTCCCCT
19951 GGGGCTGTGG CCCACAACTC TGGAGGGGGC GGCGCTTTCT TGACTCGGGT
20001 GTGATGAGTC CATCCCTTTT TCACTGTAGA AACAGCAGTC TTGGTGGTGA
20051 GCAGCACAAG GTAGGGTCCT TCCCAGGCTG GCTCGAGTTT TCCTTCTTTC
20101 CACCCTTTGA TAAGAACGTG ATCTTCAGGC TGGTGTTGGT TTACCGGAAA
20151 TTCTAGGGGT GGTACCTGTG CTAAAAGACT TTTAGTTTTG AGGGAAAGGA
20201 AAATGGAAGA TAAACCAAGT ATATAATTTC TAAGAAATGG ACCTTTTGTT
20251 TTAAATGTGG GGACATCAGC AGTGGACTTT ATAGTCCTTG GTGCCTTTTT
20301 ACTGAGAAAT TTCCTTTAGC ACCTATTTTT ATTAGATTTT AGACCAAAGA
20351 AGGCCAAACA CCATTTTATA TTTAACAGTG CTTCCTGTAT GATTCTTATA
20401 CCAGATAAGC TAAGTTTCAC CTTTATATTA GCAAGTTGTT AAACTTAATT
20451 TTAATAAAAC TTTGTAGACA TATTTATCCA ATTTTTAATG TCTGACCATA
20501 ATGTATGATT CTTATAGACT CTTTTTAACC TTTTATAATT TTTGTTAAAG
20551 AGCAGGTTAG TGCTTTAAGA AATACCTGTT GTGCTTTTAT TTTAATGTCC
20601 AGTTCACAGA AAAACTGTAT GATACCCCTT AAACTTTAGC CAATATGTTT
20651 ACACACAGAA TTTCCTTTAT AATTAACATT TCAAAACTTG CTTAAACCTT
20701 TAAAACAAAA TATTTGTTTA TTTTTAAACT TTTAATGTAG GTAAAAATCC
20751 ACATTCTTAT GGCTCCTTAT AATCCTTTTA CCAAAGGCAT ATTTTACTTT
20801 CCTTATACAC CTTGCACATA AACTGTTTCT TCAATAGCTT TACATTCAGG
20851 AGGCTTAATT ACTTTTAAAT TATACAACAT TTCTTACATA AATTCCCTTT
20901 TAAAACTTTT TTTTCCTTCA CAACTTTCAC AGACAATTCT TTGACATGCC
20951 TCAACTTTCT GACTTGTTGT AAACATCCCT TTCTTTAAAC AACTAGTTAA
21001 TTTATTTTAG GACAAGAATT TACTATATAA CATTCTTTTT ACATAAATTC
21051 TCCCTCTCCT TTTTTTTTTT AAGATAATCA TTCTTCTCCA AAGCCAACTT
21101 CCTTTATGTC TGTGGACAAG ACTGTCTAAG GCCACAAGAT TTGAAGTTAG
21151 GATAATACAT GTTACACTGT TAACTTTTAG CTAAATTTAC TTTTGTTGAA
21201 AACCTCTAAG TTTGGGATTT CAATTATTCT TTGCTATTAA TAAGACCTTG
21251 TTTAGTCAAA ATTAACTCAG AATTGGTATA GATGGCTTTT TTTTATTATT
21301 ATTATTATTC TGTAAGTACT TTAAGGCTTG GCTGAGTGCA AACAGCTCTC
21351 ACGTTTGAAC AGCACCAATT ATTAGGCAGT TTTCCTAACT CTGCTTCTAC
21401 AAGTGTTTCC TTATCACTTC CTGAATACTC ATTGTGTCTT TTTCCCTCAA
21451 TCACCCGGGA GGAACCTGTC CTGAAGGGAT TTAGATCCCC TGTTAGGCAA
21501 ACCTGCTGGG TTAAGGGGAA TTTTCAGTGG TTAATGTTAA ATCATCTTTT
21551 TCTAACAGTA ATAGCCCCAT ACTTTAAGAT TTTTGAGTTA GTAAGCTACA
21601 TTTTCACTTT TTATATATTT TTTGACTTAG GGTAGTTCTG AACTGGTGAG
21651 GTGTGCTCAC AATGAGGTTT CCTCTAAAAG TTACTTTTCT ACTTCCTTCT
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FIGURE 3G

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21701 GTTAGCAAAG CAGTTGCGGC TACAGATTGA ATGTATTCAG GCCATCCGCG
21751 GGTTACTGGG TTAAGGATTT TTGATAGGAA GGCTACTGGT TGTCAGTGGC
21801 CTCAGTGCTT TCAGGCTATG CCCTTGTTTA TACTTACAAC AAGGTGGTAC
21851 TGGAGTGTTA TAGGGTCACC GAGAAGACCT TCGATTATCA GTTATAGGTT
21901 TTAAATTTAC CCTGGCTTTT TTTTTTTTAT TATTATACTT TAAGTCCTAG
21951 GGTACATGTG CACAACGTGC AGGTTTGTTA CATATTTATA CATGTGCCAC
22001 GTTGGTGTGC TGCACCTATT AACTAAGGAA TAGGGTACAC TGTTTTTTCT
22051 TTACTACTTC TATCTCTTTC TTTCCCTCTC TGACTTTCTG TCTCTTTCTT
22101 TCTGACTCCC TCTTTGTAGC TCTGCCTCTC TTTCTCTCTC TCTGCCTCTC
22151 TCCTCTCTGT CTCTCTCTTC TCTGTCTCTG TCCTGTTTCT CTCTCTCTCT
22201 TGTTTCTCTC TCCTCTGTCT CTCTCCTCTC TCCCTCTCTT CTGTCTCTCT
22251 CTCCTGTCTC TCTCTTTCTC TCTCCTCTCT CTCTCTCCCC TCTTGTCTCT
22301 CACTCCTGGC TGTCTCTCTC TCTCTCCTCT CTGTCTCTCT CTCTCCTCTC
22351 TGTGTCTCTT TGTCCTCTCT CTCTTTCTCT CTCCTCTGTC TCTTTGTCCT
22401 CTCTCTTTCT CTCTCCTGTC TCTCCTCTCT CTCTCTCCCC TCTCTCCTGT
22451 CTCTCGCTCT CCTCTGTCTC TGGCTCTGTC TCCTCTCTGG CCCTCTCTCT
22501 CTCTTCTCTG GCTCTCTCTC CTGGCTCTCT CCTCTCTGAC TCTCTCTCTC
22551 TCTCCTCTCT CTCTCCNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN
22601 NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN
22651 NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN
22701 NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN
22751 NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN
22801 NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN
22851 NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN
22901 NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN
22951 NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN
23001 NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN
23051 NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN
23101 NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN
23151 NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN
23201 NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN
23251 NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN
23301 NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNAAA
23351 AAAGGTTGCT TTACGAGGAA GTTAAATTTA AAAGTAGAAG GCAAAGAATT
23401 GAACATACTG ACATATTAAT TCTTTGAAAA GAAATTTATA ACTGATATCT
23451 AACACTGAAG GAGGCTTATG CTTAGGGTTT TATGTTAGGA GTTTGGTTTA
23501 GAGCATAGCA TCTTTTATTT AGAAGAAATC TAATTCTTAA TATGGAATTC
23551 ACAAGTAGGA TTATGAGGAA CCCTGAAAAT TATATACAAA GTTTATTTTG
23601 TGTATTTGAA TTATTTTTTC TCTTTGGAAA AGGCATGTAT TCACCAAAGG
23651 AGTCCATGCT ATCCCCCCCA AGCTAAGACT GCTTCTGCTC ATCCTCAGCG
23701 ATTCATAGTT GCCTTAGGAT ACATTTATAG GGGACCCTCA ATTTTAAAAA
23751 CTTAGCACTG AATCAGAGAG AAAACTTGAG AGGCATTTGC GAGGTTAAAT
23801 GAGAGTGACC GATGCTGTAC AAGAGTAGGT CTCGAAATGT GGTACTTTTC
23851 TTGGGTTATC TCGTCTTATT CTCATCACAA ATGGTGAAGA AATGGTCAGC
23901 CACATTAAAG AGCAGATACT GAGATTCAGC AAGTGAGAAA ACCTGTCCTG
23951 GTTCACACAG CCAGGAAGAG GCAGAGGCAG AATCCTCACC CCACTTCTGT
24001 TTGCCTCCAA AGCTCAAGGA GAGTGAGCTT TACCCTTCAT ATTTACTCAT
24051 CCTCTTACTA ATTTGACTCT TAAGATAATC CTGAGATTTA AACCAGAAAA
24101 CTATTATGAT CCCCTTATTT GAATGAGAAT ATATGTCTAA AAATGATTTT
24151 TAAAAACACT ATTAAAGGTC ACAAAGCCAG TGAATGATAA AGGGATTGGT
24201 ACCTCTGGCT CCTATAGTTA GTTCATCCTT CAAAGAACAA AAATAGCCCC
24251 CATTTATTGA GTGCCTACTA AACTCTAGAT ATGTTTTTAA TATATGCTAT
24301 CTCATTTAAT ACCTACCACA TTCCTGTAAG GTAGGTATCA TTCATTATAC
24351 CTATTTTACA GATCAGGAAA AAACAAAACA AAACAAAAAA AAACAAGACT
24401 TTCTAGGGAA AGATGCTGAA TAGAACACAT TCTTCTACAT CCATTCCTTT
24451 CTGAAGATCT TCCTAATATG ACAGGTAGGG ATTTGTCTTA AGATTTAAAC
24501 CCACAAGGTA TGAAGAGACA GGCAGAAGAG CTTCACTATC AACGTTGCAG
24551 AAACTGGAAA GGAGACAGAG AACTAGAAGC AACATAACTG AGTCCTAAGC
24601 TTCTAGAAGG GGAAGGTGAG AAATAACCAG ACCCATGCCG TAGAACCCTC
24651 CAAAGACTCG GGAATTGGCA CTGTCATGTG CCTCTAGAGC TAGAGGTGAA
24701 GGGGAAGAGC TAAAGTAAAT GACATTGTTT GGATATCTAT TTAAAAACTA
24751 GTCATGTCCC TTCTACCAAC TTGGAAAAAG ACAAAAAAAA ATTCTCCACT
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FIGURE 3H

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24801 CCATACTATG GTTTATCCTC TGAAGAAGAA GTTTTCTTAG TGGGGAAGTT
24851 GAGTGCAGAA GATGCCTTGC TGAAAATGGA GGGATCGGGT AGATAAATGC
24901 ATACTGGATA CTGGGGCACC CAGCCTCCTC TTCCCACTTG GCTCTGATAA
24951 TACTGGCAGC CAAGGACTCA CCCTCCAGTA AAGAGAACGA CAGAATATTT
25001 TCTGGAGATT TTGACCAATC CAAGAAGGAA GATTTAAAAT TATCAACATT
25051 GGAGATTTTC TAATTCAACA TCCAGGCCAC AGCTAGAAGC AACACTATAG
25101 AAGTTTATTG CTGGCAAGAG CCACATACTC AGAATGTCCA AACAGGGGTT
25151 TAGGTCTCCA CACTTAAATA TGAGCAGACA ACCAAGGATT CTCAGGCTTT
25201 TGGGGAAGCC CTCTAATATG ACTGATAGAG ACTAAAACAA ATGAACAGGG
25251 AAAAAGTTAG CAAAAAGTAT AAGAAAGGTA AGAGAAAGCT ATGAAAACCA
25301 AAAAACAAAT AACCAGACAA AAAACAAACA AACAAAATAG ATACCAAGAA
25351 AATAGCTTTT GGAGAGCAAA AATTTGCTTT GGGAAAAAAA TTACAGCATG
25401 AATGGAAAAA TCCAAAGAAG ATTTAGAAGA TATATTTAAA GAAAATTTCC
25451 AGAATAATGA GCAAACAAAG ATATAAAATA AGGGTAAATA TAAGAACATT
25501 TAACGGCCAG GTGAGGAGTT CTAGTTTCTA AATAATAGGT ATAGAAAGAG
25551 AGAAAGAGAA AATGGAAGGG GCAATAATTA TTACATATTT TAAGAAAAAG
25601 AGTCCAGAAT TGAAGAACAT AAGTTTTCAG ATTAAAGGAG CCTATTAAAT
25651 GCCCAGCACA ATGAATAAAT CATAACATAT CAAAACATTC AACACAAGTA
25701 TATAAGACTA GAAGTTTCTA GAGAAGAAAA CTGTTACATC AAAAGGATCA
25751 GGCATCAAAA TAGCTCTAGA CTTCTCAACA GCAATGTGTG AAAAGGTAGA
25801 AGATAAGAGC AAAGCCTTCA AATTCTGAAG GAAACAATTT CCAACCTAGA
25851 ATTCAATAGT CAGCCAAACT ATTAGTCAAG TGTGAATACA ATAAAAATAT
25901 TTTTCATGGA TATATAATAT TTCAAAAAAT ATATCTCCCA TGCAATCCTT
25951 CTTACAAAGC TGTTTTAAAA TGTGCTTCAG TAAAACAAGA AAGAAGGGGG
26001 CACTGCATGC AAGAGCCAGG AATCTATCCT TAAAGAGGCA TGAAGGAAAT
26051 CCCCAGGGTG ATGGTGAAGG GAATACCAGG AAGACAGCTG TGCAGGAATA
26101 GAGATAAATA GTCCAGACTG GATTATGTCT GAGGAGAGAC ATTTTCAGGA
26151 AGATGACAAT GTGCCTGATG CACCTGAGCA TTATGAAAGG GAACTAGACA
26201 ACTGGAGAAG GGTTTGGGAT TGGATTGGGA AGGAGATGTA GAAAAGTCAA
26251 CATGTGTAAA CAAGACTGTT ACTAATTCCA GGGAAAGCCA AAAATTGTGC
26301 AAGAAAAGAA AACTAATCAT AGTTTACTAC AACTCAATTG AGCCTACCAT
26351 TTCTGTATTC ATAATAATGG AAATACCGAA TATTGATCTA ATTAAAATTA
26401 TTATGCCAGA TGTATTAGAA AGATGGAGGC ATGTTGGGAT AAAACCAAAG
26451 GAGCAAGAAC ATGAGCTAAA TCCCCATCTA CCACCTTGAA TATTCAATAA
26501 CTAATGCCTA AAATGAAAAA GAAAGGACAA TAAAATTATA CTCTTTAGGG
26551 ACATGGTGGA GATCACCCAA TGCATATCTA AAGAGAGGTA AAAGTGGTTG
26601 CTCCTTGGCT GGGAGAGATT AGAAGGGGGG TAAGTAGATC ATAGGACTGC
26651 CATTTTCTCC TTTTTAAAAA ATAACAAATC TTTTAGAACT ATTTGATTAT
26701 TTAAGCTATA TAAAGATATA GATAGTTATG GACACAAAAC TTGAAAAAAT
26751 GAAAACATTA AAAAGACTGA AATAGAGCAA AATATGAATC ATGGTTATCT
26801 TTAGATGGTT TTGTTTTTCT TCTTTATACT TTGCTGTATT TTTTATACTG
26851 ATAGCATATT CGTTTATATAT ATATGTGTGT ATATATATAT TTTACAATTA
26901 TATATACAAT TTTATATATT TTATATATAT ATTTATATAT ATACTCTTCA
26951 TTGTAAACAA GAAATTGAAG TTCAGAAAAG TCAGATAAAT TTCCTAATTT
27001 CAAATATCTT GTAAATGGTA GAGCTAGGAT TCCACTGCAA GTCTGTCTGA
27051 TGTGAAGCAT TTTTATCTTT CATCAAAGCA TTCAATCTTC GTTAAAATCC
27101 GAGAGGCAAA ATTGTCATGC CTCACCATTC TCTCCCATCT CTGAAGGTCC
27151 ATAGTGCCTC TTTTGTACAC CATACAAAAT AACACTTGAT TGGTTTCATT
27201 ATTTGTTTAC TTATTTGTCT ATCTATACAT TTATTCATAT TCATCTAATT
27251 TTAGAAAGAT GAGAGAATGG ATTCCAAAGG TACATAGATT ATAGCAAAAT
27301 AAAATAAAGT TACAAAAATG AAACAAGGGA CATTTGATTA TTCAGGTTTT
27351 GTTTTGTCAG ACTGCTAAAT GAGGCACACT CAGTTTTCCT TCTCTGCTTG
27401 GGGAGGGTAA GTGTCCTGGG ACTGAGTCCC AAGCTTCTTA TGTTTTTCCA
27451 TCAGTGCCTA GGAAAGTCCT GGGTACACAG ATACTCAATG AATGTTTGTT
27501 GGTTTGACTT GCCAGCAAAG CCGTGGCTCC TAGGGAAGTG ACTTCAGCTT
27551 CTTTATCTTC TTGGTGTGAC TATCTTAAAA GGGAGTAAGT GAGCCTTTCT
27601 TTGTAACTGA CTGTATTTGA GAATGCAGCA TGACAGACAA AACATTCATC
27651 TCATTCATGG AGAATTGTAA AATCCAGCAG AAGAGCTCTC TTTTTAACCA
27701 GTGCTTACAA TTTGTCCTTT TTCACCCTTC CTTGGCAAAT CACGCAATAT
27751 TCCTTCTTAA AAATGGGTAA AGTGCCAGCC GAACTTAGAA GAGGGACTGA
27801 TTCTATCTCT ATTCTGACCA GGTATACGGT AGACTGTAAT TTAATGTCAG
27851 CACCTTTCTG TTGCCATAAT GAGGTATATT TATTTCTGTT CAAAGATCAT
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FIGURE 3I

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27901 GCAGCCCTGA CAAAGCAAAT ACCCTCTGAC TCCCACTGTT AATTATCCTT
27951 CAGTTGCTAC AGGGTTTTCA TCCATGTCCT CACTTAGGAG AGTTGGCGGT
28001 TGTGAAGCAG ATGGAGTCCA CAATCTCAGT GGCAGTTCTT AATGCTTTGA
28051 GCTCAAAGTG TGAGTAAGTC GATGAGTGAG GCTTTTAAGA TGTAAATCCA
28101 ATATCTGCAG AGAAATCTGA AGCTGTAATA TTAGAACAAC ATTCAAATGA
28151 GGACTTCATT GACTAGCTCA TTAAGAAGTC CTTTGATAAT AGCATGTTGG
28201 TAAGACTTTT CTTAGAAGGT ACATATTATA AATGATGATG TGCTAAGAAA
28251 TCAACATAAA GGAAAATAGA AAAATTTTCC CCAAATCCAT CCTTTTTCTG
28301 TAGAACTTTA ATGATGATAC CTCATTCCTT TGTAACTTAA TTTTAAAAAG
28351 TTAATTATGC ACCTACTATG ATACGTCCAA AATGTTTTTA GGTGATGTGG
28401 ATATAGCGAA GAACAAGACA CACCCAGTGT CTTCCTTCAT GGAGTCTATA
28451 TTCTTGGCAC TGTTGGTCCT GTGTGAAGTC CTAACATTAT TTTGCTTAAT
28501 GTTTTGGCAA GAGAGGCAAC ATTGGCTGGG CGTGATGGCT CATACCTGTA
28551 ATCCCAGCAC TTTGGGAGGC TGAGGTGGAT GGATCACCTG AGGTAGGGAG
28601 TTCAAGACCA GCCTGATAAC ATAGAGAAAC CCTGCCTCTC CTAAAAATAC
28651 AAAATTAGCC AGGCATGGTG GTGCGTGTCT GTAATCCCAG CTACTCTGGA
28701 GGCTGAGGCA GGAGAATCAC TTAAACCTGG GAGGCAGAGG TTGTGGTGAG
28751 CCGAGATTGT GCCATTGCAC TTGTACTCCA GCCTGGGCAA CAAGATTGAA
28801 ACTCCATCTC AAAAAAAAAA AACCAACAGG CAACATTCTG GGCTGAAACA
28851 AAGGTAATTC ATCTGGTAAC AATAGCAATA ACATAAATAG CAGTAATAAT
28901 TATACATTAT TGAGTTCCTA TTCTCTGCCA AAAATGGTTG ATAAGCACCT
28951 TTGATATGGC TTATTTTACC TAGTCCTCAT TATAACCTTA GAAGGTATAT
29001 TGTATCTGGT CAAAATTGAA AGAAGAAATT GAAACTCACA GAGGGTAAAT
29051 AATTAAAGTT CATAGCTAGT AAGTAGTACA GACAAACCCA AAAGCAGAGT
29101 TTCATGCTCA TAGTCACCAT AATGTATTCA GAAACTTTTA GGACTCATCA
29151 CAATATTAAA ATCATGGAAC TTGGAGCCAC AAAAAGTCAG ATTTAAGTCC
29201 AAACCCTGAC CCTGGGTAAT TTAACTTTTC TGGGTTTATG TAACATATCT
29251 ATAAAGTAGC AATAATAATA TTACCACCTC ATGCTGTTTT GGTAAAAAGT
29301 AAATAAGATA ATGTATATTA AGGTATTTGG ATAGTGCCTA TAGATGTATA
29351 TATGCTACTT AATAGACAGT AATGTAATTA TTAACTATGA CCTAAGATGT
29401 GGCACAGTGC AGGTAGCAGA AGTTCTATCA TTAATCATTT ACAGATACTT
29451 ATTAAATTGC TTCAAACCCA TAAGGATAGA GGCAAGATGG AGGGGGAAGT
29501 CTAAGAAATT GATTGAGTCA ACATTTATAT AAATACTTAT CTACTGAGAG
29551 CTTCTTCACC TCAGGGTTTG GGTCACTTTA AATGCATCCT CCCTGACCTC
29601 CTCTGCCTGG CTACCTTTGG AACTCCAACC CATTCTGCAA GACCCAGTTA
29651 AAATGCTGCC CATTCCTGAA GCTTTCTTAT TTTCTAAAGT AGGAAGAGAT
29701 TTCTCCCACC TTAGAACTCC TATAAACATC TGCAGACTAG TTCTAGGCAG
29751 CCTTTAACAA AATCCTCATG GGATCTTTGA AAATACAGAT TCCCAGGTCC
29801 AGCCTCCAGA GAATCTGATT CAGATAAGGC CAATGAATCT GAATTTAAAA
29851 ACATGTATTT GTGTGATTTT GATGGGTGGA CACACTTGAG AATCACGTCA
29901 GGACCATTTA TGTGGCTCTC AATTACATAT ACACTACTTT ATATTGCAGT
29951 TGTTTATTTA TGTTATATTG CAGTTATTTA TTTATGTTTC ATCTCTTTTC
30001 CTGAGAAATT ACCTTCCTGA TAATCCAATG CAGAGATAAA TTAAGAAAAT
30051 CTGTAGGAAA GAATAGATCA TCAAGTCCCT TGCAACATTC TTCTGAGGTT
30101 GTAATAATCT CCTCTAGGAT GCTTTGCTGG ATTTCCCTGG ACTAGGTTGT
30151 CTTTTCCTGC TACTTTCTCC CATTACAGGT CTCCCTACGG CAGCACTGCT
30201 TATATCACTT GGAACTTGAA TCTATTTTGG TAAAAAAAAA GTTAAAAATT
30251 AAATTATCAG AAGGATATTG GGGATGCCTG CAGAGTAATC AAAATAGGAT
30301 CTATATTGTT ATAGAGCCAG GCACATTAAT GCCATCAGCT TTAGCCCTTT
30351 ATGTTGTGAT TTTACTTTAT TCCAAATGTC AGCTTTATCC TGTTGGATGT
30401 GCTGATCTTT TTTCTCTACA TTCAGCCAGT TCCATTCTCA TGTTCTGGAA
30451 GCTTGTGACA GAGGGGGAAT ATGCATTTCA AGATCAGAAG ATCCAGAGTG
30501 AAAATGATTG GAATGGCCTG AGTCACAGTT CCAATCCTAG AACAAGGCAT
30551 CTTGCTAGGG ATGTGAGAGA TGATAAGTGA CAGATACAGT GACAGCAAGT
30601 GGTTGATGGG ATCTGAGTTG TGAGAGAGGG TCTGTGAAAA ATGAAAGACC
30651 TGCATAAGAA GAGGAGAAGC AGAAATATGA ACATTGTTGT GAGTCAGGTC
30701 TTTACCCAAC TCTGTGCTGC TTATTCTACT TTTTTGTGCA AGATTGATTA
30751 TGTGTGTTTA ATAGAATGCA GTAAAGAACA GTGTTGGAGG GCAGCTGTGG
30801 AGTCCACTTG AGTGGGACTC TACCACTCTG CCACTTACCT ACTTTGTGGC
30851 CTTGAGAAAG GTACTTAATT TCCCTGGGTT GCAGTTTGTT CACCTAAAAA
30901 CGTGGCAATA ATAGTAATAC TGTTTCAGAG TTGGCGCAAA ATTAGGATAA
30951 TATATGTAAC ATATTTAGAA TAATGATGGG TATTCCTTAT GTAAATGTTA
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FIGURE 3J

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31001 GATGTTAGCT ACTGTGAATT TTTCTGTTGT TCCACTAGAC TGTAGGACCC
31051 CTGAAGGCAG GCAACCTTGG GCTTCTTTCT CCCAGCACCT AGCACAATGG
31101 CTGTTACTTA GTAAGCAGTC AGTAATGGTG TGTTGTTGTC AGTGAACACA
31151 GACTGAGTTC AGTGAGCAAT GTCTTGGAAA GCCTCTACTG CACCTAGGAC
31201 TTTCAGCTAT ACTGAGACAG AAAAATGAAA TCCTCTCTGG ACTGGAAAGC
31251 AGAAGCCAGA CATGTAGGCA ACCAAACTGT AACTGTTTCC ATGTCGAATT
31301 GACTTTGCCT TTAGCGAATC ATAGCACTGA GGAGTGTCAC GTTTAAGCAG
31351 CAAATTTGTA TAGCAAATTA ACATGCCAAA AAAGGCATGC AAGACTTTTA
31401 CTTGATTTTT TTCCCCTCCT CTCTGGGGAA TTTATCTTAT TTGGGTCTTA
31451 TCTTGGAATT TATCTTATCT TGAACTTATT CAGACTGCAT TGGTTTAATT
31501 TGCTATCAAC TGGGGCTATA TAGTGCACTG GAATTTAATG TGTTGTATAT
31551 GTGAAATATT TACCAAATAA CCACATAACC AAGATATGGA GGACCTACTT
31601 TAAGAGGAGA TTCTTGCAAA GCACCTTAAA AGCATACACT CAATAATCAC
31651 AATGGCATGA CTGCATACAG GGAGATAATC AGTTGTTTTA ACTTTTAATT
31701 TAAGCAGTAG CAGAATGACT TTTTGGGAAC TTAGGAATTT GGAAACCTTT
31751 TTATTCTATG TATTGAATAT CAACTATGTA ATTTAGTCTA AGGTTATATG
31801 CTAGAAACAT TTCAAAAACG AAAGCAGCAG CAATGACATC AAAAATGCAT
31851 GTCAAAAGCA AATGGTTTTA AATAGAAATA CATCATTTTA ACAATCTTGA
31901 AGTTTAAAAG ATCCTATAAA AATCACAAAC CCAGAAGGAC AAACAAGAAA
31951 AGATTGATAC ATTTAACTAC ATAAAATTTA AAACTACATT ACTGAAAAAA
32001 AATCTGAGAC AGGGTCTCTG TCACCCAGGC TGGGGTGCAG TGGTGCGATC
32051 ACAGCTTACT GCAGCCTTGA CTTCCCAGGC TTAAGGGCTC ATGTAATCCT
32101 CCCATCTTAG CCTCCCAAGT AGATGGGACC ACAGGCATGC ATCACCACAC
32151 TCGACTAATG TTTAATTTTT TTGTTGTTGA GACAGTCTCC CTATGTTGCT
32201 CAGGCTGGTC TCTAACTCCT GGGCTCAAGT GATTCTCCTG CCTCAGCCTC
32251 TCAAAGTGCT AGAATTACAG GTATGAACCA CTGAGCCTGG CTTTAAAAGT
32301 TTTTAAAATC AAAAGCCAAA TGGACAACCT AGAAAAAATA CTCCTGAGAT
32351 ATGTTAAACA GAGTTAATTT ACTTGCCATT TTTAAGTGTG CTTACATATC
32401 AAAAAATCTA ATAACTCATT AAAGATATGT AAAATATATA CAAAGGCAGT
32451 TTGCTGAAAA AATACACATA TAAATATATG CAGCTTCACT CAGCATTCAA
32501 GAAATAAAGT AAATCAATAA TTCAATCTTT TTCACTTGTC AGATGAAGAA
32551 CAGTTAATGT AGTAGTGTTG GCAAGGTGGT GGACAAAAAG TTATTTTTAT
32601 ATGTTTTTGA TATCAAGAAG ATTTGATGCA ACATCTTTGA AGAGCCAGTT
32651 AATAATATCT GTAAAATTAG AAAATTAACA TATTCTTTGC CCAGCATTTC
32701 TACTTTTATC AACTTTGCTT GTAAACAGAC ACAGAAGCCC ATCAAGAATG
32751 CTCAAGGTAG TTTTGGTAAT CATAGATAAT TTTTTTTTTT TTTTGACGGT
32801 GTCTTGCTCT GTCACCCAGG CTGGAGTGCA ATGGCACAAT CTTGGCTCAC
32851 TGCAATGTCC GCCTCCTGGG TTCAAGGGTG TTGCAGGAAG TCAGGGACCC
32901 CAAACGGAGG GACTGGCTAA AACCATGGCA GAAGAACATG GACTGTGAAG
32951 ATTTCATGGA CATTTATTAG ATCCCCCAAA TTAATACTTT TATAATTTCT
33001 TATGCCTGTC TTTACTGCAA TCTCTGANNN NNNNNNNNNN NNNNNNNNNN
33051 NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN
33101 NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN
33151 NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN
33201 NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN
33251 NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN
33301 NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN
33351 NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN
33401 NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN
33451 NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN
33501 NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN
33551 NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN
33601 NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN
33651 NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN
33701 NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN
33751 NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNACTCCCT CCCCTTTTGA
33801 AAATCCCTAA TAAAAACTTG CTGGTTTTAC AGCTCGGGGG CATCACGGAA
33851 CCTACCGACA TGTGATGTCT CCCCCAGATG TCCAGCCTTA AAATTTCTCT
33901 CTTTTGTACT CTGTCCCTTT ATTTTTCAAC GCAGCTGATG CTTAGGGAAA
33951 ATAGAAAAGA ACCTACGTGA CTATCAGGGG CAGGTTCCCC GACACAAGGG
34001 ATTTTCCTCC CTCAGCCTCC TGAGTAGCTG GGATTATAGG CACACACCAC
34051 CACACCCGGC TAATTTTTGT ATTTTTAGTA GAGACTGGGT TTCACCATGT
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FIGURE 3K

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34101 TGGCCAGGGT GGTCTCAAAC TCCTGACCTC TGGTCATCCA CCCGCCTCAG
34151 CCTCCCAAAG TGCTGGGATT ACAGGCGTTA GCCACTGCAC TCAGCAATCA
34201 CAGATAATTA AACCATCTTT CAAAATCCAT CAATAAGTTA AATATTTTAT
34251 GGTACATTTA CACAATAAAA TACAAATTAG CTACTTAAAA ATAATGAGAT
34301 CTATATGTGA TGGTATGAAT GGACAGAGGC AATGTGTTAT ACAGAAAGGG
34351 TTTAACAATG TATGCTCCCA TTGGAATGAT AGTATGTTGC TATTGCTGTT
34401 AGGAAGGAGT ACATATATGC AGAGAGAATC TCTTAAAGGG TACACAAGGA
34451 TTTGTTAATA ATGGTTGCTT CATGGAACTG GAACTGCAAA CTTGGAAGAG
34501 GAAGATAGCT TAGTTTTCAC TGAATAATTG TTGTACTTAA AAAAATTTGT
34551 AATTTTTAGT TTTCAGTGGA CCAGATATTT TGCTTCTGGT TTATAATGTC
34601 TCATCTTCAA AGTCAGCTGA GTTAGGTTTA ATTAGCTCCA TTTTACAGAC
34651 AGAGACATTG TTATTTGAAA GATTGAGTAA CTAGTCTAAG GTTACACAGC
34701 TGGTGTCCTC GTTGCCTGTT CAGTAGAAAG GTTTACATAA ACAGCAAGGT
34751 GTGCTGTTCT CAATAGACTC ACTTATGTTC ATGATTTGGT ACTTGCTCAA
34801 GCTGGAATCA ATTTTTAGAA AAAATAAAAT CTTTTGCAAA GATTTTTACC
34851 TCAAAAATAG AAAAAAAGGG CATTCCTGCC TTACCTTCTA CAAGGGTCTT
34901 CTCTGAAATT CCAAGCATCA GGGTGTTATA ACAGACTCTA AAAAGGGTTT
34951 CCTTTTTTCT TTCCTTTAAC ATTGCTTATT GCACAGCATA TTGAGACAGA
35001 GAAGATGGTA AGTGAAATAA AACAAAGGAA ATAAAAGTA TCATCACTGG
35051 GTTTCAGAAT CAGCATGGTT TATGCTAAGG GAAAGACTTG GAAACCTTGA
35101 TTCAACATAT AATTCTAAAA AGAGACAGGA AGAAATCCCA CCTTGTTTCC
35151 TCTGATTCTA CCTTTGGGAT GGGTAGGTAT GTTATACAAT AAGAATAACA
35201 TTGAGATGAC TGCTATAAAA ATAGTGGTTA AGAGCCTGGG TCCAGAATGA
35251 GAAAGGTGGA TATTGAATTT ACCTGAGTGC AACTAGGCAG ACTCAAGTGA
35301 GTTGATTTTA CCCACTCCTC CACTCAAATA CTGGGTATGG CTTTGCAAAA
35351 ACATTCAACC AGTTATCCAC ATAGTTGGTC TTAACTTTCC ATGTGACTAT
35401 AATGAATATA AACTTGCTAA TGAGCAGAGT GTGATTTTAG TGTTTAAACT
35451 ATTTTTTCCC GAATAATAGT TCCTAGATGC AGTTAATGAG CCTTATTGGG
35501 TACCCACACA AAGGAGATAG AATTGTCTGT TGGACTTTTT GAAAAACTTT
35551 CTTGGTTTTA AAAAAGGTAC ATTTCTAAAG GATTTTTATG TGTAGTTTTG
35601 ACTAAACAAG TCTTTGCCTT ACTTTCTGTT TTTAAAATCT AACCTCAACA
35651 TTAATATGTC ACTATACTGG TTATAACCAT AACAAATTAT TTCATCTCTC
35701 TGAGCCTGAG TACCCTCAAC TGTATACACT ATAAGGATGT GAAGATAGAA
35751 AGTGACATAA AAATGAAACA TGTACTGACC ACCCTCATAA ACAGATCCCT
35801 CATACATATA GAATGTCTGT GCCTGGTTGA TTAGTGAAGG AATGTGTACT
35851 CACCCAAAAG AAAAACTCTG AAATAAGTAC TTTTAGATAT TTACTTTTTC
35901 AATATTCCAA GTAATTATCA CAACATTAAG GTGCATTCAG CTTTGTGTGT
35951 TAACGTGGTA TACCTCCAGG CAACTTTTAG GATACTGTAC AGATACAATG
36001 GCTGTGAAGG CTGGGATGAA AAGACCTGTG CGAAGCAGGA CTGAGGCACT
36051 TAAGGAAGGC CTCAGAGTTA CATCTCCTTT GCCTGTTTTC TTGCAGGCCA
36101 CATACCCTAG CCCAGCCCTG TCAGCATGAG TGAGAACCAG GCTCTGCCTT
36151 TGCCCACACT AAACCACTAC CTTCAAGGCC CCACAAAGAC CCAGTGTCTC
36201 CAGACGGTCT TTCTGTCTTC TTAACACTCA GAGCTCCATG AACCAGAATG
36251 AAAGTTTTGG AACATGATCC AAGTAAAAGA CTCAAGAAGT AAACACCACT
36301 AAGGTTAACT TTGCTTTAGA GGTTAGAGAA AACACTGCAA GGACACCACA
36351 CCAGAGACTA TGAAAACCCC AAATGTATTG AAATGATGCT GATTCCATTT
36401 ACCTCCATAT TGCCTGATAA TACCCAGGTG CTACCATGGC AGCTTAAGGT
36451 GGTATTTGCT GGGAGCTATG ATACTCTTTA AGAAGTAATA GCACTACTAG
36501 TAAAAGCAGT TAGTTCCAGG CAATATTCTA TGCACATGAC CCATTTCATC
36551 TTCTTATAAA CCTCATGAAG AATATATTAT TTTCATCCTC ATTTTATAGA
36601 TGCAGAAAGG GAAGCATAGA CGTAAATTTC CAAGATTACA CAGCTATTTA
36651 TTGTTGGAAC TGAGATTTGA ATTCAGGTTG TCTGTCTTCA GGGACTGTGC
36701 TCTTAATCTC AGTGGTCATC AAACTTTTCT GTAAAGAGCC ATCCAGTAAA
36751 TATTGTGGGT TTATATACAT TCTCTATTGC ATATCCATTG GTTTTCAAAA
36801 ATAATCCTAT ACAAATTCAA AAACCATTCT TAGCTCATAG ACTACACAAA
36851 AACAGATTGC AAGTCCAGTT TGGCATTTAC TGTTCCTATT GATCAAGGGT
36901 TTAAGAACAT AGTGAGTACA CTATTCCACA TTCCCCTTAG GCAAATCCTG
36951 TATGTTTATA GTACTGTTAG ATTTCTGTTG ACAAAATAAT CCACAATTCT
37001 GACTTCATCT CTCTCTCTCT CTCTCTTTCT GATTTTGTTT GAATTTATGA
37051 GGTTTAGTTG CATTTTCAAG TTAGTCTTCC TGCTAACGAG TGATTCTTTT
37101 GTTGAACATT TAAAAAGGGA CTGTCAGGAT TGAATAAGAG AACCTCTTCC
37151 AGTCACTTTT TTTTTTGAGA AAGGATCTCA CCTGTTGCCC AGGCTGGTGT
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FIGURE 3L

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37201  GCAGTGGTGC  AATCACAGAT  AACTGCAGCC  TCAACCTCTT  AGGTTCAAGT
37251  TCCCCCTGCC  TCAATTTCTG  AGTAGCTGGG  ACTACAGATG  TGCACCACCA
37301  TGCCTAGCAA  ATTTTTAATT  TTTTGTAGAG  ATGGGGCCTC  ACTACATTAC
37351  CCAAGCTAGT  CTTGAACTCC  TGGGCTCAAG  CAATGCTCCT  GCCTCGGCCT
37401  CCCAAAGTGC  TGGGATTACA  GGTGTGAGTG  ACTGCATCCA  GCCTCTTATA
37451  GTCACTTTTA  ATCTATCATT  GGCTTTCCCA  TTAGATTGTA  CTGTTATACA
37501  AGGAAGTGAC  TTCAGACAGT  ATGGCACTAG  ACTAGAGGCT  GTGTTTTTCT
37551  TTAATAAAGG  CATAAATGAG  ATGAATTGCT  CTAAGGCTTT  AGGCTTGTCC
37601  CTTTTCTGAG  AAGTGACCTT  TGGGAGGTCA  CATTTAGTTA  AAGCAGTTTT
37651  GCTAGTATAA  ATTTACCAGG  ATCCTGACAT  GTAATCCTGT  ATCATTTTCA
37701  GTAAGGTTAA  AATGGTATAT  GAAAGGAGGT  GGTTCACGAA  ATGGATTAAT
37751  ATCAACATGG  AACTTCATGC  TTTCTAGGTA  CCTGCTGCAT  CCTTGGAGAT
37801  TCAAAATGTC  ATCATGGCAT  TCTAGGCTAG  ACTGGCAGTG  GAGAAATCAC
37851  TGTGAGTTAT  TGGATTTGCT  CAAGATAAAA  TCTTGAATTT  GCAAATAAAT
37901  CCTGGTCAGC  TTTTTTTAAC  ACTCTTGTGG  TAAATAATAC  ACAACTCAGA
37951  TTCATGTAAT  GGGTGTAAGA  AAATCATTGC  TTTGGTTATT  TCAGTATGAA
38001  ACTCAAGAGA  AAACTTACTG  AAGTGTTTTT  AAAATTATTC  TGACCACAAC
38051  CCAAGGTAAA  ACATAAGCCA  AAAAACATAT  CATGACATAG  TAAATGAAGC
38101  CAGGATTGTA  TATATATGTC  TACTCAAGTA  TATGAAATGG  AAACAACAGT
38151  TTCAGAGGCA  GTACTATGCT  TACTACATTT  GAGGCATTTC  TGGTATTTTC
38201  TATTCTATTT  AATTAAATTT  TTAGTACTTC  TTATTTTAGC  TACATTTATT
38251  TCATAACTCA  TTAATGGGTT  TTGACTCACA  GCTCAAAAAC  ACTGCCTTAG
38301  AGAATCCAAA  TGTTCACACT  ATCCATATTT  ATAAGAAGTA  ATTGTTCTGG
38351  GGTTCTTGTG  TATTCTTATA  GCTTAGTTTG  ATTTATTTGC  TAAGACCTGG
38401  CTAAGTGAGA  ACTGCAAAGA  GTTATGCCTT  CAACTACCTA  AGCCAGGAAT
38451  TTTCTGAGGT  GGCAGGGGAA  CCAGGGTGAG  CAGAAGGACA  TATCATCCCC
38501  ACCCTCATTA  AGCTTATGCT  ATAGTGGATG  AAATAAACTC  AGAAGTCAAG
38551  GAGTTTCAGA  AGAGAAGTCA  TTCCCTTGAG  TAACTATGTT  AAGTACGTAA
38601  ACAGCTTTAG  TAGTGCTTTC  TTAGTACAAG  GTGTTTTCTT  CTGATCTAGG
38651  AGAGTCAGTC  CAATTTTTTT  CTTTTTGAGAA  AATGGAGGCT  CAAAGAGTCT
38701  GTCATTTATC  TCCAGTCTCT  TCATTATTTT  GAGTCCAAGT  ACAGGATTAT
38751  TTGTAATATA  CATGCTGCCT  CACATGACTA  AGTGGGTTTT  GTGATAGAAA
38801  GGGAATTTGG  AGTTGAGAAG  AGAAAGTGAT  GATTAAGTCA  CATCATTAAA
38851  ATGTTTGACT  CTCAGATATC  TTGGAAAGAC  TTTGAAGGCA  CTCTAGCCAA
38901  ACTTTTTCCT  TCAGAAGGAG  CTTATCTAAT  TATTCTAGAT  AATAGAGAAA
38951  AACTAGGTCT  TTTAAAGAGA  CAAATTATAT  ACCATTTAGT  GTTTCACAAT
39001  ATTTTCTGAA  TAAACTTAAA  ATCCCTTATT  TGGAATTTAA  CTCATCTAAA
39051  TCCTTATTTC  AAAAACCAGG  AAACAGAGTC  AAACATTTTC  TCAGTTATCA
39101  AGGCAGTAAA  CCAAAGATTG  TCACCTGCAC  AGGAGAATCT  ATGATTTGTT
39151  CTTCTCATCA  TTATACATTT  CACGAGCATT  GACTCAAAAA  ACCATGCTAC
39201  CTATAAACTA  ATCAACAATT  GCTTCTTCTA  GGGACTGAAA  TTTTAAAATT
39251  TCAGACGTGG  AGGATCGACT  CTACTTCAAA  GCAAAATTCA  GTGGACTTCT
39301  GCACACATAT  CCATTCTAAT  CTGTTACAAG  TCTGCACTTT  GGAGATTAGT
39351  TCATGCTACA  CACTTAGAGG  TGTAATATTT  TCCTACTTGG  GAAAATTGAA
39401  ATTACTTAGA  TACAAAAGAG  TGGTTGTAGT  AAGAAAATAG  GCAAGGAGAA
39451  CATTTTAAAG  TGCTGATCCT  CGGTAAAGCC  ATACATAGGA  TGCACCTGGG
39501  AGCAGATCTT  TCTGAAGTCA  TTCTGTGCTC  AGAGATGTTT  CTCCTTACCT
39551  TGCTGCCTAT  GTCAAATTCT  CTGTGATATG  TTCTTAGAGC  CCCATGACCT
39601  CTCTTCTTAA  CTTGCAGTGG  GAGCTTGAAT  TTTCCATTTA  TTTTTGTGAC
39651  CATTTAGTCT  ATAAGAGTCT  CCGTCTTTAC  AGGGCCCTCA  CCTGACTACA
39701  GACTCCATAA  AGGCAGAGAT  TCTATTTTTA  CTCTATTATT  ACTGTATTCC
39751  CAGCACTAAG  CACTAGGATT  AATACATAGT  AAGTGTTCAA  CAGATGTTTA
39801  CTGGATGATT  AGATTGGCAT  TTTAAGGTAG  TCTGAGATCA  CGTTTTAGAC
39851  AAGATACTTC  AGTTTAGTCC  AATCTTTATT  ATTTATTAGC  TACTAAAGAG
39901  AAATTGATAA  TTACTCATGA  TATTCTTCTT  TTTTGTTTTA  CAGTCAACTT
39951  TGACCACTTT  GAAATTTTGC  GAGCCATTGG  GAAAGGCAGT  TTTGGGAAGG
40001  TGAGAACAAA  TTGAAATGAT  TAACCACCAG  CAGGGTTATG  TAGCCCAGGG
40051  AACAGAGGGT  CCAGAAATGT  TCACATTATT  GAGTTGCTGG  GACCACAAGG
40101  AAAGATAATT  AAGTGAAAAT  GTTTTTGTAA  TGGATTTTTA  TAAAATTGTC
40151  ACCACAGTTT  AAGAAAAGCG  TGTGACAGGC  AGCTACATAA  TGAACATATA
40201  CTGTTGTCAG  AATAATCTCA  TTAAACTCAA  ATCTGTTTAC  TCTCAGTAAA
40251  CTTTAAGGCT  TTTCTCTCTA  CCCTAAAGGA  GATGAAGATT  TCAGAATCAT
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FIGURE 3M

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40301 TTTCAGATTC TACCAGCTGT ATGCCCAGTA ATAGTTATCT TGTTTATGGA
40351 AGAGTTACTT ATTTTCATGT GGGAAAGAAG TCATCCGATT TCTATTTGTT
40401 TCCTCATTTG TCTAATGTTT TTATCTTAAG AAAAATACAT ATTCAGTTTA
40451 ATTTTTTTTG CAAGAAACTT CTGTATTCAA ACCCTGATTA CTAGTTTCTC
40501 AATGGAGACG TACTTTAAGA GAATAATATT TCATATAAAA CTTGCATTTT
40551 AAAATCATTT TCTGTTTACT TTTTCAGGCA TTATACAGAC CTCTAAAGAA
40601 ATTTCAAAAA CATGGACATC ATATTTAGTG TTTTTCCAGT CCTTAAAGTC
40651 CTTTTTGGTT ATATCATGTA TGGGTTGTAA ACAGAAATTC TTTGCACAGT
40701 ATTATTCAGC TTGACAGTTC AGTCATGTCT ATTTCAGTCA CTCAAAGCAG
40751 GATTAAGGAT GTTACTTGTT ATTGGAATAT TCCTGACATG GAGGCAGCTA
40801 TTTTCACCAA AATGCTGTCT TAAAAGCCCA AAAAGCAATA CCAGGCAAAA
40851 TTGTTTGAGA AAAAAGAGAT CCAAGAATTG AACTGGTGCA TAGAAAAGAA
40901 AATGAAATTT TTAATCTAAA ATCAGAGCTA AGTGGGAGCT TTTAACATCA
40951 TATAATTTGC AAATGTTAAG GATCCAAGCC ACAGCAAAGA ACATGTCTTG
41001 TTCTGTCTCT CATCACCATG ATCCATTATC TCCCTAATCA CTCTCTCACT
41051 CGGGTTTTCA CCATTAGGTC TGCATTGTAC AGAAGAATGA TACCAAGAAG
41101 ATGTACGCAA TGAAGTACAT GAATAAACAA AAGTGCGTGG AGCGCAATGA
41151 AGTGAGAAAT GTCTTCAAGG AACTCCAGAT CATGCAGGGT CTGGAGCACC
41201 CTTTCCTGGT TAATTTGTGG TGAGTAATTT TACTGGACCT CTGAATAGAG
41251 ACACTCCTGT TATCGGTGGG CTAGGGGAGG TCCCCAAATG CCTCTGGGAC
41301 CTCAGCCCTG GCTGGTATCC AGGCTCTTGA CACAATTGCA AGAAAGAGTT
41351 CAAGGATGAG TTGGAAAACA GTGAAAGTAC AGAGATTTAT TGCAAAGTGG
41401 AAAAGTACAC ACTCAAGAGA GGGGAGCATG GGTGAACTCC AGCGAATGTC
41451 ATGTAAGGGG GGGTTTGAGG CTGCTGCCAT AATGGGTTTC TTTAACCAAG
41501 GGGTGAAACA TTCATGATGA TTCCTGAAAA AAGATGGAGA TTTCTTGGAA
41551 CTGTGGTGCC AGCTATTTTT ACACCAAATA TGAATGTTCC TGGAACTGTC
41601 ATGGTGCTGG TGGGTGTATG ATTTAGTATG TTAATGAGTG TATGATGAGG
41651 TCCTAGGTGA AACCTAGGTC AAATCCAGCA CAATGGAGAG GACCCACAGA
41701 CTCTCTGAAG GAAACGACTG CTCCTGCAGG ACCCAGGCAA CTCCCCCAAA
41751 ACTGTGAGTA CCCCAACTGT GGAGGTGGGA AAGAGAGACC CTCCTCTCCC
41801 AAACACACAC CCCCACTGGA GAAGCTGAAG GTCTGTTTGC TGGAGAAGTT
41851 TCTGACTTTA CCTGGAGCTG AGTGGACTTG AAGAGCCCAG TGAAATACAC
41901 GGGGAGAAGA AGCAGCAGAA AGGCCCTGGG AGCTTGCTGG GTCCACAAGC
41951 AGGCCATTCC TGCCTGGCAC CACAGGGATC CAATGGGAGA GGAGCGGGGG
42001 TAAAATTCCA TAGGGAGAAG CAAATCTCTA GCTGAACTTG GTGACAATTT
42051 GAACAGGGTG AGAAAGCGCC TGGCCAGAAC TCAGGAGAGG GCACAAATCC
42101 AGTGTGCAGA CTCCGGGGGC AGGGGATAAA CCAAGCTCTT TTATTTCCCA
42151 GCTGGGAGCG GGGAGCCTGG GGCAGGTTTT CAAGCAGGTA TTGCTTCTCT
42201 ACTTAGAAAC AACCTGGGAG CTGTGTTGGC GGGGGAGGGG GGTTGGGGAT
42251 GGGGGAGGGG GGTGGTGGAA AGCACGGTGG GAGTGAGACC GGCCCTTCGG
42301 TTTTCATGGG AGCTGGGTGA GGCCTGTGAC TGCCAGCTTT TCCCCACTTC
42351 CTGACAATCT GCATGTTTCT GCAGAGACAG CCATAATCCT CCTAGGTACA
42401 CAACTCCAGT GACCTGGGAA TCCCACCCCC ATTCCCCACA GCAGCAGCAG
42451 CAGCAAGGCC CACCCAAAGG AGTCTGAGCT CAGAGACACC TAGCCCTGCC
42501 CCCACCTGAT GGTCCTTCCT ACTCACTCTG GTATCGGAAA ACAAAGGGCA
42551 TATAATCTTG GGAGTTCTAG GGCCCTGCCC ACTGCCAGTT TCTCCCCATA
42601 ATACCAAAGC TGATGCTCTC TGGAAAAGCA CCACCTCCTG GCAGGAGGAC
42651 AACAGCACAA AAATAGAATA TTAACCAAAG CTAAGAACCC TTACAGAGTC
42701 CATTGTACTC CCTGCCACCT CCACCAGAAT AGGCACTGGT ATCCACAGCT
42751 GAGAGACTCA TAGATGGTTC ACATCACAGG ACTCTGTGCA GACGACTTCC
42801 AGTACCAGCC TGGAGCTGGG TAGGCTAGCT GGGTGGCTAG ACCCAGAATA
42851 GAGATAACAA TCACTGCAGT TCAGCTCACA AGAAACCATA TCCATAGGAA
42901 AGGAGGAGAG TACTACATCA AAGGAACACC CAGTGGGACG AAAGAGTCTG
42951 AACAAGACTT TCCCTCTGAA AGAGCCTACC CAAGTGAGAA GGAACCAGTA
43001 ATATGACAAA ACAAGGCTCT TGATGCCCCC CAAAAATCAC ACTAGTTCAC
43051 CAGCAATGGA TCCAAACCAA GAAGAAATCC CTGATTTACC TGAAAAAGAA
43101 TTCAGGAGGT TAGCTATTAA GCTAATCAGG GAGGAACCAG AGAAAGGTGA
43151 AGCTCAGTGC AAGGGAATCC AAAATATGAT ACAAGAAGTG AAGGGAGAAA
43201 TATTCAAGCA AATAGATAGC TTAAAGAAAA AACAATACAA AATTCAGGAA
43251 ACTTTAGACA CACTTTAAAA ATTGCAAAAT GCTCTAGAAA GTGTCAGCAA
43301 TAGAATTGAA CAAGTAGAAG AAAGAAATTC AGAGCTCGAA GACAAAGTCT
43351 TCAAATTAAC CCAATCAAAC AAAGACAAAG CAAAAAGAAT AAGAAAATAT
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FIGURE 3N

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43401 AAACAAAACT CCCAAGAAGT CTGATATTAT GTTAAATGAC CAAACCTAAG
43451 AATAATGGGT GTCCCTGAGG AAGAAGAGAA TTTTAAAAGC TTGGAAAACA
43501 TATCTGAGGG AATAATTGAG GAAAACTTCC CCGGCCTTGC TAGAAATCTA
43551 GACATCCAAA TACAAGAAGC ACAAAAAACA CCTGGGTAAT TCATCGCAAA
43601 AAGGTATTTG CTTAGGCACA CTGTCATCAG ATTATCCAAA GTTAAGATGA
43651 AGGAAAGAAT CTTAAGAGAT ATGAGACAGA AGCACCAGGA AACCTACAAA
43701 GGAAAACCTA TTAGATTAAC AGCAGATTTC TCAGCAGAAA CCCTACAAGC
43751 TAGAAGGGAT TGGAGCCCTA TCTCTGGCCT CCTCAAAACA ATTATTAGCC
43801 AAGAATTTTG TATCCAGTGA AACTAAGCAT CATATATGAA GGAAAGATAC
43851 AGTCATTTTC AGACAAACAA ATGCTGAGAG AAATTGCCAT TACCAAGTCA
43901 CCACTACAAG AACCGCTAAA AGGAGCTCTA AATCTTAAAA CAAATCCTGG
43951 AAACACATCA AAATGGAACC TCTTTAAAGC ATAAATCACA GAGGATCTAC
44001 AAAATAAAAA TACAAGTTAA AAAGCAAAAA CAAAACCAAA AAAATCTGCA
44051 GGACCCAGGA GACCACCCCC AAAAAAATGT GAGTGCTCCA ACTGTGGAAG
44101 TAGGAAAGGA AGAGCATCCT TTCCTGAACA CACACCCCCA CTGGAGAAGC
44151 TGAAGGTCTG TTTGTGGGAA GAACAGCTTT AGCTCTTTTT TGGTTTTTTG
44201 GAAAAAAACC CAAAGTACAC AGGCAACAAA GAGCATGATG AATGCCAACG
44251 GTACCCTCAC ATTTCAATAC TAACATTGGA ATGTAAATGG CCTAAATGCT
44301 CCACTTAAAA GATACAGAAT CACAGAATGG ATAAGAACTC ACCAACCTAC
44351 TATGTGCTGC CTTCAGGAGA CTCACCTAGT ACATAAGTAC TCACATAAAC
44401 ATAAAGTAAA GGTGTGGGGA AAGGAATTTC ATGCAAATGG ACACCAAAAG
44451 CGAGGAGGGG TAGCTATTCT TATATCAGAC AAAACAAACT TTAAAGTAAC
44501 AGCAGTTAAA AGAGAGACAA AGAGGGACAT TATATAATGG TAAAAGGCCT
44551 TGTTCAACAG GAAAATGTCA CAATCCTAAA CATATAAGCA CCTAACACTG
44601 GAGCTCCCAA ATTTATAAAA CAATTACTAA TTGACCTAAG AAATGAGACA
44651 GACAGCAACA CAATAATAGT GAAGGATTTT AATACTCCAC TGACAGCACT
44701 AGACAGGTCA TCAAGAGAGA AAGTCAACAA AGAAACAATG GATTTAAACT
44751 ATACCTTGAA ACAAATGGAT TTAACAGATA TATACAGAAC ATTTCATCCA
44801 ACAACTGCAG AATACACATT CTATTCAACA GAGCATGGAA GTTTCTCCAA
44851 GATAGACCAT ATGATAGGCC ATATAATGAG CCTCAATAAA TTTAAGAATA
44901 TTCATATTAT ATCAACATTC TCTCAGACCA CAGTGGAATA AAACTGGAAA
44951 TGAACTCCAA AAGGAAACTT CAAAACCATG CAAATACATG GAAATTAAAT
45001 AACCTGCTCC TGAATGGCAT TGGGTCAAAA ACAAAATCAA GATGAAAATT
45051 TAAAAATTCT TCAAACTGAA TGACAATAAT GACACAACCT ATCAAAACCT
45101 CTAGGATACA GCAAAGGCGG TGCTAAAAGC AAAGTTGATA GCCCTAAACG
45151 CCCACATTGA AAAGACTGAA AGAGCACAAA CTGACACTCT AAGGTCACAC
45201 CTGAAGGGAC TAGAGAAACA AGAATAAACC AAACCCAAAC CCGGCAGAAG
45251 AAAGGAAATA ACCAAGATCA AAGCAGAACT AAATGAAATT GAAACAAAAA
45301 AAAAAAAAGA AAGATAAATA AAACAAAAAG ATGGTTCTTT GAAAAGATAA
45351 ACAAAATTGG TAGACTATTG GCAAGATTAA CCAAGAAAAC AAGGGAGAAA
45401 ATCTAAATAA CCTCACAAAG AAATGAAACA AGAGATATTA CAACTGACAC
45451 CACTGAAATA CAAAAGATCA TTCAAGGCTA CTATGAACAC CTTTATGCAC
45501 ATAAACTAGA AAACCTAGAA GATATGGATA AATTCCTGGA AAAATATAAC
45551 TCTCCTAGCT TAAATCAGGA AGAATTAAAT ACCCTGAACA GATCAATAGC
45601 AAGCAGCGAG ATTGAAACGG TAATTTAAAA ATTACCAAGA AAAATGCCCA
45651 GGACCAGATG GATTCACAGC AGAATTATAT CAGACATTCA AAGAAGAATT
45701 GGTACCAATT CTTTTGACAC TAAGGAAACC TCCCCTAATT CATCCTATGA
45751 AGCCAGCATC ACCCTAATAC CAAAACCATG AAAGAACATA ACCTAAAAAG
45801 AAAACTGCAG ACCAATATCA TTGATGAACA CAGATGCTGA AATCCTTAAC
45851 AAAATACTAG CTAACTGAAT CCAACAGCAT ATCAAAAAGA TAATCCACCA
45901 TGATCAAGTG GGTTTCATAT CAGGGATGCA GGAATGGCTT AACATACACA
45951 AGTCAATAAA TGTGACACAC CACATAAACA GAATTTTTTA AAAAATCACA
46001 TGATCATCTC AGTAGGTGCA GAAAAAGCAT TCAACAAAAT CCAGCATCCT
46051 TTTATGATTA AAACCCTCAG CAAAATCAGC ATACAAGGGA CATAGGCCTT
46101 AATGTAATAA AAGCCATCTA TGACAAACCC ACAGCCAACA TAAAACTGAA
46151 CACATTCCCT CTGAGAACCA GAATGAGACA AGTATGCCCA CTCTCACTGC
46201 TCCTCTTCAA TGTAGTACTG GAAGTCCTAG CCAGAGCAAT AAGACAAGAG
46251 AAAGAAATAA AGGTCATCTA AATCAGTAAA GAGGAAGTCA AACTGTCACT
46301 GCTTATTGGC GATATGATCG TTTAACTTGA AAACCCTAAG GACTCTTCCA
46351 GAAAGCTCCT AGAACTGATA AAAGAATTCA GCAAAGTTTC CGGATACAAG
46401 ATTAATGTAC ACAAATCAGT AGCTCTCCTA TACACCAACA GCAACCAAGT
46451 AGAGAACCAA ATCAAGAACT CAATCCCTTT TACAATAGCT GCAAAAAAAA
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FIGURE 3O

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46501 CAAAACAAAA CAAGACAAAA CAAAAAAACA AAAAAAAACA AATACTTAGG
46551 AATATACTTA ACCAAGGAGT AGAAAGACCT CTACAAGGGA AAATTACAAA
46601 ACACTGCTGG AAGGAATCAT AGATGACACA AACAAATGGA AACATGTCCC
46651 ATGCTCATGG ATGAGTAAAA TCAGTATTGT GAAAAATAAC CATACTGCCA
46701 AAAGCAATCT ATAAATTCAA TGCAATTTCC ATCAAAATAC CACCATCATT
46751 CTTCACAGAA TTAGAAAAAA CAATTCTAAA ATTCATATGG AACCAAAAAA
46801 GAACCTGCAT AGCCAAAGCA AGACTAAGCA AAAAGATCAA ATCTGGAGGC
46851 ATCACACTAC CTGATTTCAA ACTATACCAT AAGCCCACAG TCACCAAAAC
46901 AGCATGGTAC TGGTACAAAA ATAGGCACAT AGACCAATGG AACAGAATAG
46951 AGAACACAGA AATAAACTCA AATACTTACA GCCAACTGAT CTTTGATAAA
47001 GCAAATGAAA ACATAAAGTG GGAAAAGGAC ACCCTTTTCA ACAAATGGTG
47051 CTGGGATAAT TGAATAGCCA CAAGTAGGAG AATGAAACTG GATCGTCATC
47101 TCTCACCTTA TACAAAAATC AACTGAAGAT GGATTAAGGA CTTAAACCTA
47151 AGACCTGAAA CTATAAAAAT TCTAGAAGAT AACATTGGAA AAACCCTTCT
47201 AGACATTGGC TTAAGCAAGG GTTTCATGAC CAAGAACCCA AAAGCAAATG
47251 CAATAAAAAC AAAGATAAAT TGCTGGTACC TAATTAAACT AAAAGAGCTT
47301 TGCATGGCAA ACGGAAGTCA GCAAACAGCC CACAGAGTGG AAGAAAATCT
47351 TCACAATCTA TACATCTGAC AAAGGATGAA TATCCAGAAT CCTACAATGA
47401 ACTCAAGTAA ATCAGTAAGG AAAAAACAAT CCTATCAAAA AGTGGGCTAA
47451 GGACATGAAT AGACAGTTCT CAAAAGAAGA TATACAAATG GCCAGCAAAC
47501 ATATGAAAAA ATGCTCAACA TCACTAATGA TCAGGGAAAT GCAAATCAAA
47551 ACCATAATGT GATTCCACCT TACTCCTGCA AGAATGGTTA TAATAAAAAA
47601 AAAATCAAAA AACAGCAGAT GTTGGCATGG ATGCAGTGAA CAGGGAACAC
47651 TTTCTACACT GCTGGTGGGA ATGTAAACTA GTACAGCCAC TATTGAAAAC
47701 AGTGTGGAAA TTACTTAAAG AACTAAAAGT AGAACTACCA TTTGATCCAG
47751 CAATCCCTCT ACTGGGTATC TACTCAGAGG AAAATAAGTC ATTATTCAAA
47801 AAAGATACTT ACACATGCAT GTTTACAGAG CACAGAGTTG CAACCCAAAT
47851 GCCCATCAAT CAATGAGTGG ATAAAGAAAC TGTGGTATAT GTATACATGA
47901 TGGAATACTA TGCAGCCATA AAAAGGAATG AACTAACAGC ATTTGCAGTG
47951 ACCTGGATGA GATTGGAGAC TATTATTCTA AGTGACGTAA TTCAGGAATT
48001 GAAAACCAAA CATCATATGT TCTCACTGAT ATGTGGAAGC TAAGCTATGA
48051 GGATGCAAAG CAATGAGAAT GATACAATGG ACTTTGGAGA CTTAGGGGGA
48101 AGAGTGGGAG GGGGGCGAGG GATACAAGAC TACAAATGTG GTGTAGTGTA
48151 TACTGCTCAG GTGATGGGTG CAACAAAATC TCACAATCAC CACTAAAGAA
48201 CTTACCCATG TAACCAAAAC CACCTTTACC CCAATAACTT ATGGAAAAAT
48251 AATCCAGCAC CACATTAGGT TTAGTCGGAC TTAGCCAGCT TGGCTTACAC
48301 CCTGGTTTTT CAGGTTCTTA TCATTCCCAG TTTATGCAGC TGTTTCAACA
48351 TTTTCCTTTT GCTAGTCATG TGAAACTGCT GTCTGGAATT TTCTTTTCTC
48401 CTGCTACCAC CCTTTATTAT TCCTGTCTCA CTTTCATCTT CATCCCTACT
48451 GTTACATAAA TGCATCTTGA TTTCTAGGCA AGCATTTGTC AAATTCTCAT
48501 TAGGATCTTC CTCAGGGTCT TTTGTTCTCC TTAGTTTCTT TGGCTTTATA
48551 GTGAAAGAAC ATTTTTCTTT TATTGTCACT AACAAATACT TCTTGGTCAG
48601 TTGTCACAGT TCCCCTTGTC CTTGAGGTCA ATATATATAT ATTTTTAAAC
48651 ATTGTAATTA AATATGCTGA CTGGGAAGGA GTTCAGATGT CTTACTAGTT
48701 ATTAGATACT TTCTTTCCCC ATGAACTGCA CGGGAGGAAC TTTGGTTACA
48751 AAGCTTGGCC TCATCAGCTG ACTTGAGGTT GATATTTAGA ATTTATACGA
48801 AGCACTTTCT CCCTTAAAAT AACTGGCAAT AAAACTGTTG CTTTGTAGCG
48851 TATTTCTTAG GCAGCCACAT ATATACCTGT AAGTTAGACA AGGATAGGTG
48901 CTTCCTTTGT CAACAAATAG CTTTTGCAGA GCTGAAGCTA ACTTGTATCA
48951 ATGACTAGAC ATTAAGTGAC TGTGATCTGC GCTCCAAGCT ATTTCCATAA
49001 TCCAAGGCAT AGAAAATGGC AGAGAAGCTT GCAGTATCTG TTACCTCCTG
49051 TTCTTTTCTT GTGTGTCAAG GTCTTTGTGT GTCACCTTCC ATTTTATTTT
49101 ACATTTTAAT GCGTCCATTA TGTTAAGTGG TGTTTCTTAA AGCTAATTCA
49151 GGATGACTGT TATTTAAATA TGCATACCAA GAAGTTCTGA CTTACCAGCA
49201 AAGAAAAAAA AGGGTCTTTA TTCAGAGAAT GCTAATGGAA AAATAATTGA
49251 GGTTTTACTC TGTGTTTAGG GACATCCTTC TGGAGAAATC AGTACATAAA
49301 ACCTGCCTCC ATCCATCTTT AATTATTACA GTTCATTTAA TATACAATTT
49351 GCTCAAAGCC TCTATGCCAC AGTTGAAAAG AAGATGGTTT TATGTGACTT
49401 GGAAATAGGT CTATTACAGT TTATGCACTA CTCGGATATG GTAGAGTCTA
49451 ATTTCAGCTT AAGCTCAGTG TATTTAATCA GTATCTTAGA GTGGCCTATT
49501 CAAAATGCTG CCATGTAAAA AGCTAAAATG GATGCAGCTC TTTCTTCCCT
49551 ACCCTTAGCA ATCATCAAAT TGCCTTTCTT CCCCTCTCTC TGCATCCTGA
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FIGURE 3P

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49601 GAATGACAAG ATACTGTCAC TTCACAACCT CCCTTTGTTC AAAGTCACAT
49651 TTTTCTTCTT AAAAAGTTTA ACCAACTAAT TTTTTTTTTT TTAAGACCAG
49701 GGACCCATGA TAAGGCCTTA GCATTTTACC TTCTCATATT TGTCTTTCAT
49751 CGCTGTGTGG GCAAAGTTGA TTTCATTCTG TTCCTTTTTT TAAGAAAATG
49801 GGTATTGTGA GGCTTTAAGC TGGCCAAAGA TGATAGATTT TGCTGTTTGC
49851 TAATTTGGTG TCATTCCAGA CAACATTCTG TTCTCCATGC ATACTGACCT
49901 GGTGATAACA TGACATATAA CCTATTCTTT CCTTCTCACT TCTCACATTG
49951 AACCTCACAG TGGAACACTA GGCATCATTA ACAATGATAG AAGAAAGAGA
50001 GGAGACTTAC CTCCACCCAG TGATTCTGGT ACTACATTCA AAACTAGAAA
50051 CTAACTGGGA GGGGGAATTC TTAAAGTACA ACAGCAACTC CCTTTGTCTT
50101 CCAAACCATG AGAAAAATCT TCACAAATCT GTATCATTCT TCCTAATAAA
50151 TGCTTTTTGT TTTAGTAAGT ACAATATATT CAATGTAAGT TTATCTTTCC
50201 ACATTTATAA ACCATCTTGC AGTGCTTTTG AAGGTGTGAT TGTGAGTGTA
50251 TTAGTCAGTT CTCACATTGC TATAAAGAAA TACCTGAGAC TGGGTAATTT
50301 TTAAAGAAAA GAAGTTTAAG TGGCTCATGG TTCTGCAGGC TGTGCAGGAA
50351 GCATAGTGGC TTCTGCTTTG GGGAGGACTC AGGAAGCTTC CAATCATTGT
50401 GGAAGGCAAA AAGGGGAGCA GGGCATCTCA CATGGTGGGA GCAGGAGCAA
50451 GAGAGAGGAG GAGAGAGTCA CTACACACTT TTAAATGACC AGCTCTCTTA
50501 AGACCTCTAT CACGAGAACA GCACCAAGAG GATGGTGTGA AACCATTCAT
50551 GAGGATCCAC CCCCATGATC CAATCACCTC CCACCAGGGG CCACCTCCAG
50601 CATTGGGGAT TACAATTCAA CATGAGATTT GGGTGGGGAT AGAGATGCAA
50651 ACCATATCAG TGAGTAATTT ACTTCATCAT TTTTAAGTCA CATGGTTATA
50701 AGATAGGGTT AATGTGTGTA ACTTTACATT TATAAATGAA ATGAATAAAG
50751 TGCTATGGCC AGTACCCAGC ACATAGTAAC AGGTGTCTTA CAAATATTCG
50801 TTCTTTCCTT CCTTACTTCA TGAAGTTATG ACATTCTGAA CTTGCCCATC
50851 TCCTATGGTT CATTGTGGAC ATCCAAAGGA CAAATCTAAA TGGTGCTTGG
50901 CCCCAGGACA TCATGGAAAG CTGTATGTGC AGTGTCAAGG GGGTTATCTT
50951 CAACTCATTC TCTATAAGAG CATATGTTGC TTGTTTGTT TTGTTTTCTA
51001 TCCTCATTCT GCAANNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN
51051 NNNNNNNNNN NNNNTATCTT TACCACCCGA AAGCATTAAA AGCTTAAGAA
51101 GCATTGTATT ATTTATAAAG TAACAGCAAT ACTTTTAAAA TTTCTGCCTT
51151 CTTTGTGTAC TCTATTTTAT GGATATGCTG TGTAGGCCTC TCAATAATAC
51201 TTTCAATAAT CTCATTCATG CTAAAATGCC CCTAGCTTCT GGAGATTTAT
51251 AAAATTCTAG TTTTCAGGCT GAGGGTAAAC AAATTGTTCC TTTTTTAAGT
51301 GAGTTAAGAT TAAAAAGTTT GTGTGTGTGA ATAGGTATAA ATGTATACAT
51351 ACATATGCAT ATATTTATAC ATTTATACCT ATACACACAC AAACATATGT
51401 TTTTCAATAT CATATATATG TATTATATAT ATGAATATCA CATATATATG
51451 TGTGTGTATA TATATATGTG ATATTTGAAA ACTCTTCTGC TCATTGCAGT
51501 CAACTTGAAA AACAGAAAAT TACCTAGAAA AATGAAAATT CTTCAATAAT
51551 TCTTATCACC TTGTAACAAT CACTTCTAAC ATGTTGGTGT ATGCTTTTCA
51601 GTAAAATGTC TGCATTTGAT TTTCTGTTTT TGATCATGCA TTAGCCTCAA
51651 TCATTCCTTC TTTCACCTAT ATGTTTACTG AGCATCGAAA ACAAGTTATA
51701 ATTTGATTGC TAGGTGAAAT ACAAGATGCA CAGTTATATT TGAATTTCAG
51751 ATAAACAACC AATAATTTTT AGTATATGGC CCAAATATCA CGAGATATAT
51801 TTTTACTGAA AATTTTTATT TATCTGAATC TGAAGTTTAA TTATGCATGT
51851 TGTACTTTTA TTGGTTAAAT CTGGCAATCT TAACTGTAGT AGAAACCCAT
51901 GCTATGGGGA TATCTTGGTG AGCAGAAATA GACGGAGCCC TGATATTAAT
51951 CAAATGGCAC ATGAATATAT AAACTGTGAG AAGTATATAA AGAGAGTAAG
52001 TATGAAGAAC TGTGTGTGTG GTGTGTGGGT ATGTATGTGT TGGGGTTTCA
52051 GAGAAGAAG GTAAGTAGTC TGGGGGCAGG GACGTTAAGG AGGAAAGAAC
52101 ATTTGGAAAT AAAATTCAAC CTGACTTGCC TCCAGGGACC TGGCTACACT
52151 CAGGAACAGT CTTCAAATGT AGGCCATGTT ATCAAGTGAA TGCTGCCAGA
52201 CAGGGCTGGC ATCCAGGAAA AGTAAATAAA ATCTTCTTGT GCGTCTGTCT
52251 CTGAGGGCTC TTCACAAAGC CCTGGCAACC CACAGCCTGA AAACAAATAG
52301 GCCCCAGTCT TTCCCAGCAT AGTTGATTCC CCAGGTGGCT TTTGTTAATT
52351 GAGATTAAAC CTGTAGCTGC ACACAACTCC TCAGGGCCTC TATCTCTTTA
52401 CTCATGTCTT TGTCCCTGTG GATAGAAGGG GTCCACATGT GGTTTCAGGA
52451 AATTAGGACA CCAGATCATC TGTTTTAACT GGAAAGAACT ACCTGTACTG
52501 AGAGTGTGAC AAGGTCCTTT CAGACTCTGA ACATAGCCCA ATAAATGGTA
52551 TCAACCTTAA ATAACGAGAT TCTGAAAATA TGATTAAGTA TCGAGTTTGC
52601 TGGAGCCCAG AGCTTGAGGA TGCCCACCTG GGAGCACAGA TTCACTTTGC
52651 CCAGAATGTA CACTCCAATT AGCAGCAGTT ATAAGTGGGG TTTTAAGAAA
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FIGURE 3Q

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52701 AAAAGACAAG GCAGTTCCTA AGTTATTTAC CAAAAATTTA CATTAAAATA
52751 ATGTAAGCTA TTGATGGACT ATGCATTATT CTTTATATCA CAAATTACAG
52801 GAACACAAAG ATAATGGGTG AGGCAGCTAG TCAGGAACAA AATGGCTTTA
52851 AAATACTGTC CTTGAGCATG GGTTTGAGGC TGTGACTGAC ATCCCATACT
52901 CATGTTTCTC TAAACCTAAT AAATTGTGCA TATCTCATAT AGCTCAGACT
52951 GCTCTGAGCT ATTTTTGTTT TCTCATTTCC CCCCTTTTCA TCAAGATTTT
53001 GCAAAGAAAG CATTGTGGAT GAACTTAAGC AGTTTTGGCT CCTTTTATGT
53051 TCAGGAACTT AGTCCTGCAT TGCTAGGAAG TCTTATTCCC AGATGGTCCT
53101 GTCCCACATT TGGGGGAAGG GGAAAGGATG AGTCTTAGTG GGGATTTTAA
53151 CACCATCAGA AGCAAAATTG GGATGGCATC GCAGGGTGCC ACAAATGAGA
53201 CCTCACCCAA GTCACTAATT TATGTAGCTA CTGTTGCTTG TGGGATCATC
53251 TCCAGGCTTC AGAATACCAT GCAGTTAGTT TTCTCGGAAT AAGTAAAACA
53301 ATGAGCTATA CATAGTAGAA ATATAATACA CATAACAATT ACAATTAAAA
53351 AAAAAAAAGA ATTTCTATGC CTGAATGAAA AAAATATCTA TTCCATTGGA
53401 AAGTCAACTA AAAACATCAT GAAGAAAATT AAAATCCAGT CCTTTCTTAG
53451 AGACTTGTTG TAGCAGGAAA TAATTCAAGA TTTAGATCAA ATTGTAGGAA
53501 AATAATAAAA ACTAGAAAAC AATGGTCAGG GCTGAATTTA AAAACAGGTG
53551 TGCTATAATT TTCTTCTGAA CCATAATTTC TCTCTCTTCA GTTCACTATT
53601 TCTACCCAAG ATAAATGTTA TCAGGACCAA CATACTTGTA AAATAAGCTT
53651 TAGTATTATA TTTGGCCTAA TTATTTGCAT TAAGTGCAAC AAAAATAATG
53701 AATGGCCATG TACGCATTTT TAAGTTGGCT TTGCTGGAAC TTTTTCATAA
53751 GGAATCTCAG ATTAGACTTT TAAAAGCCTC TCTAAACTAG ATATTGAAGC
53801 CAATAATTCA CCATCAAACT GCCTGTAGCA TCTACATAAA TTGGGTGAAT
53851 TTCTCCCTTC TTCAGGTTCT GAAATATATT GAGGTTTCTA GGCCTGTCAA
53901 ATGATGACAT TCTTTACTTA CTGCAAGGTC AAAAAACTTG TGAGGGTACC
53951 ATGTAGACAA GGTATCAGGT CAGTTTTCCA AAAGGACTAT TGATTTGGCT
54001 CTATAAAGTC AACTTCAATT CATCAAAGCA GTTTGGTCAT ATCTGAAAGT
54051 ATGTCATTTC ACCCAAAGCC TTGGTAAAAT GACCAGCCTT AGTAAAATGA
54101 CCAGTGTCTC CAACTGTGTA CTGTTACAGA AGAAAACAGG TTCTTACTGA
54151 ACTTACACAA ATAACAATAT TGCCATAAAT AAAGAGTATT CACAAATAGT
54201 TTCCAAATTC TGGAGGAATC AGGTAGAGAG TAAGATGTTT CAATTTTGCT
54251 CATAAAAGTA TACTTTACTT AATTGTTGTA AGCTCTAAAT AGCTCAAAAA
54301 AAATTCTTGA CTTTGGAAAA CAAAACAAAA AGAATCAGCA ATGTTCCAAA
54351 CAAAAAAAGT CATTAAAAAA ATTTCAGTCC TGGCCAGGTG CAGTGGCTGA
54401 TGCCTATAAT CCCAGCATTT TGGGAGGCCA AGGCAGGTGG ATCACCTGAG
54451 GTCGGGAGTT CAAGACCAGC CTGACCAACA TGGAGAAACC CTCTAAAAAT
54501 ACAAAATTAG CCGACGTGGG TGGCACATGC CTGTAATCCC AGCTACTCGG
54551 GAGGCTGAGG CAGAAGAATT GCTTGAACCT GGGAGGTGGA GGTTGCGTTG
54601 AGCTGAGATC ACATCATTGC ACTCCAGCCT GGGCAACAAG AGTGAAACTT
54651 CATCTCAAAA AAAAAAAGAA AAATTTAGTT CTCTATCAGT TCAGTTCCAT
54701 GTAGTTAACT CTTGTTCTGT TTGATATTGG GTTAGCAATC TTCACGAACT
54751 GATGAACTTT TATATTAGAA TTCTGAAAGT TTTTACATAA TCCATTGATA
54801 TGATTTCCAA AACCTTCAGA AACTTGTATT CGAGAGTACT TCTCAGAATC
54851 CTTTTCATGA ATTTCCTTGA AGGATAAGCA AATTTTGGAC TGTAGCTGAT
54901 TATAAACCAC TTTTTATGAA GAATCTAAGT AAAATAATAA TTGTCTGTAG
54951 ATGACAAAG ACTTAAAGCA GTCTTAGTTA AAGACACAAT TGACCAGGAA
55001 ATTTGGTTAT GCCTGTAGCA TACAACAACT TGACATAACA ATCGTAATTA
55051 TTACTGATCA TATATACCAA AACATATTGG AACTTTTGGA ATCTCATTCA
55101 ATTTTGGAAC AGATATTAAT CATATTAATA CATTTATACA AATATATTCA
55151 AAGAAAGTTA AACATCATTT CTTATTTGAC AATGCTTTCT GTATGATTTA
55201 AACATATCAA ATAAGCCTGA TCTGCCTCTC TGTAACTTCT AGGGGACCTC
55251 ATATCTGAAA AGTTATTTCG AGGTAAAAAA AAAAAAAAAA AAAAAAAGGA
55301 CTAAATTTTA ATTTGAAATA TGATTTTGGA AAGTTTGTCA AATATCAAAG
55351 GTTTAAAAAA CTTACTCAAA ATATTTTTAC AGGTCACTGT AAAATAATAG
55401 TCATTTATTT AGCCAAAGTG ATAATTCCAA GATTTCAAAA GCAAAAACTT
55451 TTACTATTTG GTAGAAAGGA GACTGCGTTC CCAATCAAGA GACCTAATAG
55501 GGACAGCATG AGGCAAACTC TTCCCTCCTT TTTATAAGGA ATCTCAGATT
55551 TTACCTTAAA AAGCCTCTCA AGGCTAGGTA TCTTTGAGAG GTTACCTTTT
55601 TTTTTTTCTG TTTTTCTTTT TGAAGTTTAA TCAAAGGCA AACAAATCTT
55651 TTACTGTCTC TTATTAATAC TATATAAAAT TCTTATTCAA AGGAGAATGC
55701 CAAATTTATA TTAGTGTGTT GTCAATACTA AAGCTAATTT TAATTAAACA
55751 TTATAAACAA ATCCATACAA TCTCAGTCAG CTTTGACTGC AGAAGATAAG
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FIGURE 3R

55801 ATTTTCATAA ATCTTTTATA ACCTATTACA ATTTTCTATT AAAGAGAAGA
55851 TCAATGTTTC AAGAAAACCC TGTGGTTCCA AAAGAGGGGC CCAGACTCTG
55901 GCCTTGCACC AGTGAGCTTT TGAGATTAAT GTTCACTTTT TAGAAAAACT
55951 TATAAACAAT TCTCTTCTAA TTTTAGCCAA CTTGATCACA CACAAAATTC
56001 CTTTCACAAG ATTAATCTTC CATAAACCCA CAACTTGCTT AAACCTTCAG
56051 TTTTGTCCTA TACTTCTTTT ATTTTGAGAC GGAGTCTCAC TCTGCCCAGC
56101 CTGGAGTGCA GTGGCATGAT CTCGGCTCGC TGCAACCTCC GCCTCCTGGG
56151 TTCAAGCAAT TCTTCTGCCT CAGCCTCCCG AGTAGCTGAA ACTACAGGCA
56201 TGCACCACCA TGCCGGGCTA ATTTTTGTAT TTTTAGTACA GACGGGGTTT
56251 CACCATATTG GCCAGGCTGG TATACTTCTT TTTTAGATTG GCATTCTATC
56301 TTAGGACAAA ATCTACTTTC CTTTCTCCCT TATCATTTTG ACCACACAAT
56351 GCTCTCTTTC ATGCAAATGA AAAATTACTG TCATTTCAAC TCCCTTTACC
56401 AAAAACACAT CTTAATTTCT TTATATACCT TATGTATAGA ATTGTCTCTC
56451 TTATATCTAG TCATTTTTTT TTTCTTTTTT CTTTTTTTCT TTTTGAGATG
56501 GAGTCTCACT CTGTCGCACA GACTGGAGTG CAATGGTGCG ATCTTGGCTC
56551 ACTGCAACCT CTGCCTCCTG GGTTCAAGCA ATTCTCTTGC TTCAGCCTCC
56601 CAAGTAGCTG GGACTACAGG CATGTGCCAC CACACCTGGC TATTTTTTTG
56651 TATTTTTAGG AGAGACAGGG TTTCACTGTG TTTGCCAGGG TGGTCTCGAT
56701 CTCCTGACCG CATGATCTGC CCGCCTCGGC CTCCCAAAGT GCTGGGATAA
56751 CAGGCATGAG CCACCGCGTC TGGCCATATC TAGTCATTTA AATTACATAC
56801 GATAACTACA ATTTTAACTC TTAGGAACGC TAATTTACAG TGAAATCTGA
56851 GGAAGTAATT TTGAGCTGTT TTATGCCAGT ATTTATAGAT GAAAACCATT
56901 TCATAATTTT TATAAAGTTG TTTCCTCAAT TATTTTGTTT ATTAACAGAT
56951 CTAAATATAT TTAGCTTTTC TACACCATAT AACTCAGACA TTTTATGGTT
57001 ACACAATGCT TAATTTAACA TGACTTTACG ATTTAGTTAC TGAAAAAGAT
57051 TTTTGAAACT GAAAAGTTCA TTTATACACT TCTATCTCAT TTACATTCAT
57101 TTAATTTAGT TTATTCATTC TTAACAATTA TGCTTGAATA GTTCATTAAA
57151 CAAAAGTAGC CACCATCAAG TTATTTCTTT GTTAATCATT TTTATAGCCT
57201 GCAAATGTCA GGCAGTTGCC ACCTAAGCAA GAACCCGAAA GCTAAAACAG
57251 AGATATTTTG CTGATCAGAA GGCACGGTGG CTTTCATTAA ACCAACAGTA
57301 TTAACTGGTC TTATTTACCG AAGATTTACC CAAGTTATGT GAACTAAAAG
57351 GGATTTGAGT TATTTTCTAT TTTTCTGATA AAATATTTAA GTGTTTCCTT
57401 TCTCTTTTGG CCAATTAGAA CTCATTCATA TATTTTTGTA ATAAATTTTA
57451 CATACACATG ACACATATAA ACATGCAGAC ACACACAGGC AGATTTTATA
57501 GCTTTGTAAG TTTCTTCATT TGCCAGTTTT CAATAGTTTC TCTCCCACCT
57551 TTAGACTGTC AAGCCCTAAA CAATTGTTAG CTAGGCAACC TTAAATTTGT
57601 ACTTCTAAAG GGATGACTCT TAGCTGAAAC AAAGTAAAAA AAAATAAAAA
57651 TTACACTTCA AAAACACAGA GCGGAGCTCA AACTAAGGGA GCAGGTGTAT
57701 ATAGGTAAAG GTCCAGTTAA GACAAGATGG CCAAGGAAAG CATCTTAAGT
57751 AAAGGTAGGA CTTGTATAGA TTTAAACCAA TGTTAAATTT CTCATGACTC
57801 AGCTCTCCCT CTCCTCCAGG TGCACAGAGG CAGAAACCCT TACAAATGGA
57851 GATTTCCTTT ATCAATGTAA ATTTCAATAT AGCCAGCTAA ATGCCAGCAA
57901 GGTATATTTT GGAGAACTGT TAGAGGCAGT GAATCTGTAT GTGTCTGCAG
57951 CAACTTCAAT TCTTGCCTAC TCTCAAAATA AAAAATTCAA CTGAGGGGCA
58001 TAAGGTAGAA TGAAAGACAG AGGCAATTTT TAGAGCAAAA GGGAAAGTTT
58051 ATTTTAAAAG TTTTAGAGCA GGAATTAAAG GAAGTAAAGT ACACTTGGAA
58101 GAGGGCCAGA TGGGCAGCTT GAGAGATTCA AGCACACGGT TTGACCTTTG
58151 ACTTGGAGTT TTATATGTTG GCAGGCTTCT CGGGGGTTGT TGCTTCTCCC
58201 CTGATTCTTC CTTTGGGGTG GACTGTCCGC ATGTGCAGCA GCCTGCCGGC
58251 ACTTGGGAGA GGCCGCATGT GCAGTGTGTT TACTGAAGTT ATGTGCATGC
58301 TTACTTGAGG CATCTTTTTT TCCTTACCAG TTGACTGTTC CTAGAGGAAG
58351 GTCATATACC AGTTAAACTC TACCATTTTT GCCTCTTAGT GTGCATGCTT
58401 GAGCCTACTC GCCCACCTCC TGAGATCTTA TCAGGAACCT ACTGATCATC
58451 AGTTTCAGGG TTTTTCTATC TACTGGGAGA TTGCCTTTTC CTGGCGCCGG
58501 CTGCAACCAA ATATTATTTG AGAGAGACAG TTTAACAACC ACCTGACCAT
58551 CACCTAATGG TTGTCTGACA TTCCTTGGTG GAGGTTGGGG GTGATCTCCT
58601 GCCTTGCCCA TGTCTGCCTG CCTACTGTAA CAGACCAACT TAGTTAAATA
58651 GGTGGGCTTT TCAACTTAGT TTGTTTCTTG GTGAGATGAC TGACATCATT
58701 GTGAAGCTCT TTAATGAACA GGGCAAAGAA AGCCTTCTCT ATGCCTGGAC
58751 TCGGCATGGA CAGCTCTGGG AAAGAAGAAA GCCTATTTTA CCTGAGGGCC
58801 TATCTTTTAT AAATATTTTG TTCAAATTCT TTCTTTTAAA ACAAAGGTTC
58851 TTTTTCAATG ACTTACCAAA CCAATACACC TTAACCAAGG TTATGTCTAA

FIGURE 3S

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58901 ACCAAGGATC AACTAGGCAT TTCCAAAGAG TGGCAAAGTA GTCCTCACAA
58951 GATCCAGAAC CAAAGACAGC TCAAAGAAAC AAATGTCTTG CTCACTGCAA
59001 ATAGAATACA ACCCATATTT CTGTCCAGCC GTATTTTCAA GGATCTCAGC
59051 TTCTCTGTTG AGCACCTACT CACGGAGGCC CCAAAGCCCT ATATGCCCCA
59101 CAGATAGAGA CAGGAAATCA AAAGCTGTCT CTGGAAGGGA AAAGAATCAA
59151 TAACAAATGG GTACCTCAGA AGGTCAAGAG TTATACAAAT GATTTTAAAC
59201 AAATAGGACT GCTTTCCTGA CTGGGAATCA AACCTGGGCT GCAGTCATGA
59251 AAGCAGAATC TTAGCTGGTA GACCACAGAG TGGAGTGCTT TTTTGTAAAT
59301 CCTTCAGGAG ATCCAAGCAG GCAGTTTGAG CATATAAAGG ATTTCAACTC
59351 ATTTCAGATC TGATCACAGC TGGAATGCTG TTTAGCTAAT TTCCTGCATG
59401 TTAATATTTC AAAGATATGA TGAGATTTGT ATCTGCAAGG GATTGTGAAG
59451 TCCAGCAGGG CATTTGAAGG ATATTGTCTG GGCCGGGCAT GGTGACTTAA
59501 ATGTGCTGGC TTAAAATCCC AGCACTTTGG GAGGCCAAGG CGGGTGAATC
59551 ACTTGAGGTC AGGAGTTTGA GACCAGTCTG GTTCACATGG TGAAATCCCA
59601 TCTCTACTAA AAAATACAAA AAATTAGCTG AATGTGGTGG CACGTGCCTG
59651 TAATCTCAGC TACTCAGGAG GCTTAGGCAG GAGAATTGCT TGAACCTGGG
59701 AGGTAGAGGC TGTAGTGAGC TGAGATCACA CCACTGCACT CTATCCTGGT
59751 GACAGAGCAA GACTCTGTCT CAAAAAAAAA AAAAAAAATA CTATCTGATG
59801 TTGGGTCAAG AAATCATCAG TGTCATTCAT TAGACCTGGT ATAGACAAAA
59851 GTTTGTTGGA TCTGTATTTT TATAATCTCT GTAGTATCAT TCTTGTTCTG
59901 TAGTTGTTTC ATTTGTTCTC TCTGTTTAAA AATTATCTTC CTAGGAGATG
59951 GATGGGAGCT GAGGGAATGA GCAGAAAGGG ATGAGTTTAG ATCACAGGAG
60001 TAGGAGGAGA TGGAGCAGTT AGAGGTGAAA GAGAAAACCT CCAAAATCTT
60051 ATTAAATTTA GAAATAGTTT CAAACATACT TTTGTTCACC TCTTGAATGG
60101 AGGCAATTTT TTCTTTTAGG ATTTCTTTTA GAAACTTGTA GGTACTATTG
60151 GAAGTAAGTC TCTCACTCAA TTTGGTTCTA AAACTAGCTT TTTCTAATTG
60201 TGTGTGCAAA CAAACTAATT TAGGTATTTT AAAAGGTACC ACATTTTGGC
60251 CATTGTCAGT TGGAATCATT CTGAGTTATG CTCTACTAGT TTTCTAAATA
60301 TTTGCATGAA GAGGCATGGT AAGTATTCAG TATGAATCGA GCTGGCATTT
60351 CTAATGGTGG ATCTCTTCTT AAGGAGGAAA CCTCAGTTTT AGATAGTTGA
60401 ACTGCCTTCA GAATCTGGCC AGTTTTAAAA ACTACAGTTG TTTTTTCTTA
60451 AGCCACAAAG ATTTACTTAT TTTTCAAGAG AAACTATATT CTTCTTGGCC
60501 AAATTTTGTA TTAGAGGAAA GGTTACAAAC TCTAATGAAT AAGACAAAGA
60551 AAACCTTAAC TTCAGAGAAA AGTGAAAATC ACAAAACAAA GTAAATATAA
60601 TCTCTAGAGA ATAACACATG AAACTCCTGT CTTTCAGTAG AGTTTCAATT
60651 CCAATCCCGC AGAGTTAAGA ATGTGTATGG CTTGAATAAA GTCTGAATCC
60701 TCAACTAACC TGGGAGTATT TGGATACCGA GATGGCTGCC AGATCTGGTG
60751 AGGTTGGGTG AACCAAGCTG TTGATTCTGG TACTGTTACA GGAAAGCAGT
60801 CCTGATCCAT ACCCCAAGAG AGGGTTCTTG GATCTCACGC AAGAAAGAAT
60851 TCAGGGCAAG TTTGCAGAGT AAGGTGAAAG CAAGTTTATT AAGAAAGTAA
60901 AGGAACAAAA GAATGGCTAC TCCATAGACA GAGCAGCCCT GAGGACTGCT
60951 GGTTGATCAT TTTTATGGTT TTTTTAATAA TATGCCAAAC AAGGGGTGGA
61001 TTATTCCCTT CCCTTTTTAG ATCATATAGG GTAACTTCCT GACATTGCCA
61051 TGGCATTTGT AAACTGTCAT GGTGCTGGTG GGAGTGTAGC ATTGAGGACG
61101 ACCAGAGATC ACTCTCATCG TCATCTTGGT TTTGGCCGGC TTCTTTGCCG
61151 CAACTTGTTT TATCAGGAAG GTCTTCATGA CCCGTATCTT GTGCTGACCT
61201 CCTATCTCAT CCTGTGACTT AGAATGCCTT AACTGTCTGG AAATGCAGCT
61251 CAGTAGGTTT CAGCCTCATT TTACCCAGCT CCTATTTAAG ATGGAGTTGC
61301 TCTGGTTCAC ACGCCTCTGA CAGTACCAAC ATTCCAATTG TCACGAACTT
61351 GAGGGGATCA CTGAAGCTCC ACTTTAGATC CCATCTGGGG TGGTAAAATG
61401 TCAACGTGAA ACAAGATTCA GAAAATATGA TTAAGTATAG CATTTATTGG
61451 GGCTCAAAGC TTGAAAATTG TTATCCGGGA GCATAGATTC AAGTTGCCCT
61501 GAATATACTC CAATTAACAG CAGCGACAAG TGGGTTTCTA CGGAAAAAAG
61551 AAGAGGCAGT TTCTAACTTG TTCGCCAAAA ATTTACGTTA AAGTAACGTA
61601 AGCTATTGAT AGGCTACACG TTATTCTTTG TATCACAAAT TCCAGGATCA
61651 CGATGATAAT GAGCCAGGCA GCTAGTCAGA AACAAAATCC CAGGCATCAG
61701 TGTGGGGATA TGACTGAAGT CCCATACTCC TGTCTCTCTG GGCCTGACAC
61751 ATTTTGCATA GTTCATATAG CTCAGCCTTC TCTGAGCTAT TTCTCTCTTC
61801 TCAGTGGCTT TCCTGGAAGC AGCCTCCATC ATATGTGACT CAGAGTGCTA
61851 GCATTTCTTC ATGGGTTTAT AAACCATAAG AACTCAAGGT GGCCTTCAGA
61901 GCCACAGCAT CAACAATATT AACTTCCCTA TTAGTAGTGT TCTATTACTT
61951 TGGGTTTTAC ATATATTATC TCATTTATTC ATCATAACAA CCTGGTTGAT
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FIGURE 3T

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62001 AGGGATTATT ATTCCCATTC TATTCCTGAA GAAACTGAGG CTCAAAGGAG
62051 CTAAAATATT TTCCTATAGT CACACAGCTA GGAAGTGGCA GAGCGAGGAC
62101 TCAAACCCAA GAATCCTGAC TTCAAAGCCT CTGCTCTTCC TGCTGCACTA
62151 TACCATCCCT ATACACATCT CTGAGACTCC TGTAAAAATA TGTAAGGAAC
62201 AGGATTTATT TCATTTATTG TCTTTCATAT CCCACAAGAA TACAAACTGT
62251 GTAAGGCAGG TATGTCTGTA TGTTTTTTAT CACTGCCTCA TTCCCCATCT
62301 TCCACAACAG TGCCTACCGC ACAGTAAGTG CTCGATAAAT ATCTTTTAAA
62351 TGAGCATGTG AATGAATGTG TGTTAGTGTT AGGGCTAAGG CCTTTGGCTT
62401 CTGGTTAATT GCCCTTTTTG CCATTATGCC AATGTCATTT GCACACTCAC
62451 AAACATACCC TCATATAATC ATATGCACTT CAGTTTCTTT GCAGGTCCTG
62501 GGTTCAGACA AATCTGAGTT TGAATTTCTG TTCCACCACT GGGTAACTGA
62551 GTGAATTTGG TCAGTTATGT TTGGTATTTT ACTTAGTTTC CTCACCTGTA
62601 ATTAGGAATA ACAGGAATAC TCATGTCAGT ACTACTTTGA ATGACAGTGA
62651 TAAGAATATG TACTTCAAGC ACCTCACAAA GTACGTGGTT GATAAATGGT
62701 GACTTTACAC AACAACTGAG TGACACTTCT TCTGGCACAG GGGCCAAGGG
62751 AAAATTTCCC CTTCACCCTC TGAAGGTTCA CTGAGAATCA ACTGATAAAA
62801 GGCAGATTAA TAGGAGAAAA AGCACACAAA ATTTGTTTGC AATATGGAAA
62851 TTCACAGAAA GGGGTAGATG GTTGACACTT TTATGCCATC TTGAGGTTAC
62901 AGAAAGAGCT TGGAAAAATA GATTATGGGT GAAGGGAGAG AAAGAAAGTC
62951 CTGGGGCAAA GGTGGTCCTT GTTATGTAGA TGAAATCTCA CAAGTAGCAA
63001 CTCTCAGAAA GAATAGATGA TAGTCTGTGG TTGGGAGATC TGATCATGGG
63051 GAGGTCCTCA GAGAATGCCT GGTTGTTTAT TTCACTAATG TATTTTTTTT
63101 TCCTATAGAT ACAAATCATC TCCATGAAAG GTAGCTTTTC AGGGTTATTC
63151 CTGTGTGCAT GCCTTCTTCT GAAGCACCAT CTCAAGATAT GTCAAATAAG
63201 TGTATTTGGG GTGAAATATT TTTGGTTTCC TTTGCTAGAA ATGAAATGTC
63251 CCTGCTTCCC CATAGCCAGA AAAGATTCTT GAGTGGACAA CTGCACCTAA
63301 ACTTGAACCT GAGCACTAGA AAGTCTTTTG TTTTATTCTA TGTTTTTATA
63351 AATTTAAATC TAATTTTTTG AATATAAAAT AATACATATT TTGTAAATGT
63401 GGAAACACAG AAAGTTCTAA TGAAAAAATA AAAACCTGTA TTTCATCACG
63451 CAGAAATATC TGCTGTATTA GTTTTCCGTT GCTGCGGTAA CAAATTGCCA
63501 CAAACCTGGT GGCTTGAGAC ATCATAGATT TAGTATCTTA CAATTCTGGA
63551 AGTCAGAAGT CCAAAATCAG TCTCCCTAGG CTAAAATCAA TGTGTCACCA
63601 GGGCTGTGTT TCTTCCAGAG CCTCCAGGTG AGAATCTGTT TCATTATCTT
63651 TTCTAGCTTC TTGAGGCTGC CTGTATTCTC GGCTTGTGGC CCCTTCCTTT
63701 ATCTTCAAAG CCAGCAGCAT ACTATCTTCA AACCTCTCTC TGACTCTGAC
63751 TTCATGTTCT CCTTATTCAT CTTTTAAGGC CCCTTGTGAT TACATTGGGC
63801 CTACTTGGAT AATGCAGGAT CACCTCTCTA TCTGATGATG GGCCTTAAAG
63851 TCCCTTTTGC CACAAAAGAA AACATATTTG CAGGTTCTGG AGATTATAAT
63901 GTGGACAGCT TTGGGGAGCC TTTATTCTGC TTATTACAAA CACTATTAGT
63951 ATTTAGTGCA ATTCATTCCC ATTGTTTTCC CTATATTTTT CAACATATTT
64001 CACTTTTTAC TATCTATGCC ATTCACAAGA TTGCTTATTT CAAGCAACGT
64051 TTTATTGTAA TTGTTTTCTG TTATCAACAT AAAGTAATCA AAAGGGTCAG
64101 AATCTAGTTT AAAGTGAGTT TATTCGAGTA CAAAGTTTGA GGACAAGCCC
64151 CCCAGGAAAC AGAATTCAAG GAATGGAAGT CAGAGTTCCA AAGTGTAGAC
64201 ATTGGGGATC ATTTATAGAC AAAGTTCAGG GAAGTTTAAC AGAATTTCAC
64251 CATCTTTCTA TGTAAGGTTT AATGCATAGT TACAACAATC TGATTAGTCA
64301 AAGTGGTCTT TTTCTTTTGA GAAATGTATA TTTAAACATT CTACTCTGAA
64351 GATGTAATTG TCATGGGGCC TTGGGCACCA TCATGTCTGA GTTAGGTACA
64401 AGACTATAGG GAGGCAGTTA ATCTATAACA AAGATCAGTG ATTGGAAAGG
64451 GGAGGTCTGG TCTCTTCTAG TCATTTATAG AATAAGAACA ATGAGGAAGA
64501 GAGGTAAGCT ATAATCTAAG ATGCAGAATT GCAGACATGC CATGCGACTC
64551 ACTCAGTTTC CAGGGCTTAA CTTCCCCCTT GTCAAAATCA ATTTAGAAGA
64601 TCCTGAAATT TTATTTTATT TTATACTTAT ATTATTAAAC ATGTTTTATT
64651 AGAATGTTTC ATTGTTGTGG GGAGAATTCC TAAATTTCCT AAGCATAAAC
64701 ACTCTTTGTT TCTTTTCAGT ATATATTTCT TCCCAGTACA TGTTATTTGG
64751 ACCTAAGTCT TCTGGGATGG CAATAGAGAT GCAATGGAGG TCAAATTCCA
64801 TCCTTTTTAG AGGAATCTAT ACAAATTAGA GCTAGTAAGG ATATAAAAGA
64851 TCATTTTATC AGGTGCATCA TCCCTAAACA TACATACACA TTTACACACA
64901 TAATGTAAAA TCCTGTTAAA AGAAGACGCT TCCCAATATT CAAGGGCTGT
64951 ATAGACGTGC TTTTAGATTA AGAATTAGAT GCATTATGAC AGATTTTGCT
65001 ATGTAACAAA CTGCCCCAAA ACTTATTAAC TCAAAACAGC AAGTATTGAT
65051 GTCTCATGAT TCTGTAGATT GGCCAGGAAG TTCTTCCAGT CTGGGCTGTT
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FIGURE 3U

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65101 ATGTGAGTCA GTGATTCAAA ACTATCCATC TAGGCCTTGA AGGCGGGGGC
65151 TAGCCTAACC TTTTTCTTCT GCCATGAGAC TAACCCTGGC TTCTTCACGT
65201 GCGGGTGGAA GGGTTCCTAA CAGCAACAGC TGACAAACTT AATGAGCAAG
65251 CACTTTTTCA GCCTCTGCCA CAGTCACATT TTCTATCCTA TTGGCTAAAG
65301 TAAATCACGA AGTCAGGCTC AGATTCAAGG GGTGTAGAAA TAGGCTCCAC
65351 TTCTGATGAG TGGCACGGCA AAGTCAACAT TGCAAAAAGC CAGGCAGAGA
65401 TATTACTGTG GCCAGTTTTG CAAACAATCC ACCGTAATAC ATAAAATATG
65451 TTTAAGCAGT CCACAAAATG ATCAAGGAAA TGGTAGAAAC TATAAACACT
65501 GCAAGAACTC AGAGCCACAT GATGTTATTG AGTCCTTGTA GTGCTCTGAA
65551 AGGGTTCAAG GAAGAAGTTG TTTTGGCATA TGACCCTGAT GAACTTGCAA
65601 AAGTAGAGAA GAAGGGAGCA CAGTTTCTGA AGAAGAACTT AGTAGAGAAG
65651 TGTTATTCTG TGGCCAGTAC GCAGTAATTG TTCCACCTAG AGATGTTGAC
65701 TGACTGATGA ACAGGAAGCT GAGTCTTTAT AATGCAGATA TTCACATATT
65751 CATTTACTCA TCCTTTATTG AAAACAACGC AAGGAGCCAC TAGAAAATTT
65801 AAGCTCAAAA GAAACTCACT GGATGGATAT GGGGTAAAGA TTCAGAAGCA
65851 CAGCTGAAGT AGCAGGTTTC ACAAAGATTA GGGACAAAGG GCAATCTGGA
65901 AATCTAGGTA GCAGGAACTA TTGAATAGAC TCTTAAGCTG TCTGGGCGGA
65951 CATGAGTCAG CTCCAACCAA TTTTCTAACC TTGTGTCACC CACTCAAGAT
66001 TGAAAGTCCT GGGAGAGAAT CCAACTGGCC TTGCTCAGAA AACATTCCTG
66051 CCCCTTAGCT CAAAGAAAGA ATAAAATAAA TGACTCCTGG ATTGTTAGCC
66101 TAAGCAACTT AGATGATCAT GTCATTCATT TAGATGGGGA GATTGGAGGA
66151 GGAGCAGATT CATTGTGAAA ATCAGGAAAA CTCTTTTAGC TCTGTTAATT
66201 TTGAACTGCC CCTTAGTAAT TCAGATAGAG CTCTTGAATA GGCAGTAAGT
66251 GAATCTGGAG TTCAAAGGGA AATTCAGGGA GTATAAAGTC CAACAAAACA
66301 AAAATATGGG AATCACTGGC TGTTAGATGC CATTTAGACC AGGGACTTGA
66351 AGGGAGCACC TTGGGAAAGA GACTAGATGG AACAGAAAGT CTGAGGACTA
66401 AAGACATTGC TCTCTAATAG TTCTGGTAGA GGAGGAAGAT TCAGGAAACT
66451 AGACAGAAAG ACAACAGTCA TGAAGCTAAT CAACAAGCTA TGGGTAAGTC
66501 AGGGGAGTCT GCCATCCTGG AATCTTCCAG AGAGAAAAGT TTTTCAGAAA
66551 GGAAGGAGGG AAAACCATTT CAGATGCTGC TGCAAGGTCA AGAAGAAGAA
66601 GACAAAAAGA GCAGACCCCT TACTTGAGAA GATAAATATT GTGACCTTGT
66651 CCCAGTGTTT TGGGAGGCTG AGGCAGGAGG ATCACTTGAG GTCAGGAGTT
66701 TGAGGCCAGC CTAGGCAACA TAGTGAGAAC TCATCTCTAC AAAAATATAAG
66751 AATAAAATAA TTAGCTGAGT AATCTCAGCT TCTTTGGAGG CTGAGGTGGG
66801 AGGATCCCTT GGGCCAGGAG TTTGAAGTGA TTACTCCACT GCACTCCAGC
66851 CTGGGTGACA GGGCAAGACT CTGCTCTAAA AAACTAAAAA AAAAATTAAA
66901 AAAATATATT GAGATTGTTG CAGAACTTTC TCCTTAGGTC AGCTAAAACT
66951 GGGCTCTTGT CACATGACCA GGGAAGATTA GGCTTGCAGA CACATAGAAG
67001 GGTGAGGAAA ACATTTATTG GGAGAAAAGG AAAAAGAAAG AAAAACCCTC
67051 AGCAAAGCGA GAGGGAGTCT TGCCAACAAC CTCCTGCCTC ACAGATAGGT
67101 TACCACACGG AAACTGAAGA GGCCAGGCTC CTCCCCCTGC AAACAGCGCG
67151 AACTTCCCCT GGCTCCACCC ACTTCCCTCA GTGCGCAAGT GGGCATTATT
67201 TAGAGAGAAT GAGCCAGGAA AGCGCGGGCT TCATCCAGGA CCAGCAGTCC
67251 GGTTTTTCAG CCTTCAGGCT GTTTTAGACT TGGAGGCTGG GTTTCTCCGG
67301 GACCCTTGGC TGTCTCCTGT CTCTATCAAG ATCTTAATAA GAGCCAACTC
67351 CACATGGTGG GACAAAAGAC CAAAGGGAGT AAAGGGAGAG GCTTAATGAG
67401 AAAATGAGAA ATTAAATCAT TTAATGAGTG ATTTTATTTT CCAAGTAGAG
67451 GAGGAGAGGT ACAAAATGAG TTTTGAGATT CATGTTGTGA CAGGTAGCAA
67501 TAGTGTCTTG CCATTTCTGT ATTGTATTCC ATTGTATAAA TACTCCATGG
67551 TTCATTTACG TTTTTTACCA TTGATAGGCA TTTGGATCGT TTGCAATTTG
67601 AGACTTTCGC AGAGTACTAC TATTAACATT CTTATTTGTT CTTTTGGCAA
67651 ACTCCAAAAT ATGTGTACTT TTGTACACAT GTAAACCCTA GGACCCAGTG
67701 GAGCGTAGTA CTTGATTTTA CGNCGTGTAG ATTAGAGTGC AACAGATCTT
67751 TAGTATACTT TAGCTGAGTA GAGTAGCAGA TAATGCTGGA CGAAGACGAT
67801 TGTCGTGCTC GTGTAGTAAC CTGTTCTAGT CTTGCGTGAG AGCACCTCTC
67851 TAGCCGCTGT GACGTCGTAC CTAGTGTTCA AGTAGCTGAG GAGCAGTGTC
67901 ACAGTAGGAC GTCCGCACCA GAGTTTAGTT CGGGTCGACT ATGATGTATG
67951 TGTACTAGTA GTGTAGTATA GTAGTACACG AGTCGTAGAG GAGTAGCCTT
68001 AGAGANNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN
68051 NNNNNATTGG AATCCTCCCC TGTGCTCTAA AGATGTGTAC TTCTTATTTT
68101 TCTACACATA TTTTGGAATA TAAACTTAAG AATTAAATCA CTGGGTCCTA
68151 GGGTTTACAT AGGTTTAGCT GGCAAACAAT TTTCCAAAGA GCTTGTGCCA
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68201 GTTTATACTC ACATCCGCAA TGTATGAAAA GTCAAGTTGC TCCAAAGCAT
68251 CACCAACACT GGATATTATC AGTTTATTTA ACTCTGGGTG TTCCAGCAAA
68301 TGTGTAATGG TATCTCCCTG TGGTTTTAAT TTGCATTTTT CTGGTGACTT
68351 ATGAGTTTGG GCATATTTTT GCTTATTGAC CATTTATAAT CCCTTTGTTG
68401 GGAAGTGCTT GTTTGACTCT TTTAACCATC TTTCTATCGG TTGCCTCTTT
68451 TTCTTATTGA TCCATGAAAG CTCTTTATAT ATTCTATATA CAAGTCTTTT
68501 TAAAAGTTTT TTAAAAACTT TTATTTAGCA CATACCAAGT CAGGTGTTGT
68551 TCCAGGTGCT GAAATGGAGG AGAAGGAAAT TTTCAGAAGA TATGTGGCAA
68601 AGAGAAAAAA GCGTTAACCT TTGTGATTTG TGTTATTTGT TACTATCAAG
68651 TTGGCAATAA TAAATATTTA TTATAATTTG TAACACATAT TTAAAATGTA
68701 TTATATATAA TATTTTATAT TGTATCATAT ATAAAATCAA CAGATTTTAA
68751 TTAATTCAAA ATTCAGTATC TTCACTGACA TGTGTTAGCT TCCTAGCACT
68801 GGAATGTCAT TTGCTTGCTT ACATATAAAG GTATAATAAA ATTTTAAATC
68851 TTCTGCTCAG ATAAAGAAGT AGTGAATTAT CTAAGATGTT TGGAATGACT
68901 TAACATAAAT ATTTCTAAAG GGAAAGGGAT AAATCACATA ATTTTTCTGC
68951 ATGGAAACCA AATAAAACAA ATAAAAAGAA AGATGCGTTT ATCAGTAGGG
69001 AAAGTGTCTA GAAAAAGTAC ATATAACTAT GCCTGACAAT AGGCATATAG
69051 CCTACATGTA ATTGATACAT TTTAGAAGAA AGTGTGGAAT CATTTTTAAT
69101 ATTATGTATG TAGAACTCTA CCCTGAGTCA GGAGTTTCTT GTCATATGTT
69151 GAGGAGGGTA GAACAGAGTT ACTAACACTA AATGAGACAT TGAATAACCT
69201 ATCTTTTGTT TTTATGGGTA AAAAATATAG CGACCATAAT ATACCAGAAG
69251 TAAAAGAAAT ACAAATTAAT ATCTAATTTA TTATATATAT GGAATGAGCT
69301 GTGAAACTTC ACCAAGAAGT CTTTCTTTGG GGCATATAAA CTATTTGCAC
69351 AATCTCTGAC CTTCTTTTTC ACTGCAATAA TGGTTTTTTT TTTAACAATA
69401 AAAAATGTTT GGACTTAATG TGGTACAATT TATCAATCTT TTTCTTTATG
69451 CGTAGTGATT TCTGTGTTCT CTTTAAGAAA TTTTTGTCTG GCTGGGGACA
69501 GTGACTCACG CTTGTAATCC CAGCACTGTG GAAGGCCGAG GCAGGCAGAT
69551 CACTTGAGGC CAGGAGCTTG AGACAAGCCT GGCCAACATG GTGAAACACC
69601 ATCTCTATTA AAAATACAAA TATTAGCCGG GCGTAATGGC ACATGCCTGT
69651 AAATCCCAGC TACTTGGGAA GCTGAGGCAT GAGAATCCCA TGAATCCTAG
69701 AGGTGGAGGT TGCAGTGTGC CGAGATCATG GCGCCAATGC ACTCCAGGTT
69751 GGGCGACAGA TCCAGACGCT GTCTCAAAAA AAAAAAAAAA AAAAAAAAAA
69801 TCTTTGCCTA TGCCAACGTG GAGCTATTCT ATCCTGTTTC CTAGAAGCTT
69851 CACTGTTTTA GCTTTCACAT TTAGATCTAC AGTCTAGGAT CAAGTTTTAT
69901 TTTGTCTTCA TATAAATAAG TAATTGACCC TTAGCCATTT GTTGATAAGC
69951 TTATACTTTC CTTACGTCAC CACAGAACCA CATTTGTTAT TAATCAAGTC
70001 ACCATCTATG TATGGGTTTC CTGACTCTGT TCCATTGATT CATTTGTATA
70051 CTCTTGCATA TTTATCACTC TGTTTTAATT ACTGTAGTTT TATACTGGAT
70101 TTTCAGTAAT TCATCTTTGG ATTATGTTGG CTACAGTTGG TTCTTTAAAA
70151 TTCCATATAA ATTTCATAAG TAGCTTTTCA ATTTGTATTT TAAAGCTGCT
70201 GGTATGTATA TTGGGTACAT GGAGTCTATA GATTAATTCA GGGATAACTA
70251 ACATCTTTTT AAAATATCAA ATTTCCAATT CATACATTTT ATATATATAT
70301 ATATATGTGT GTGTACATGC ATATACATAT ATATGCGTAT ACATTTCCTT
70351 ATTTATGTAG ATATTCCTTA ATTTCTCTCT TTGGTTTTAG TTTTTCATGT
70401 AGAGGTCTAG CGTATTTGTC TTTAGACTGA TGACTAGGTA TTTGATAAGA
70451 TTACAAGTGG TATTATTTAT CAAAATTGTA TTTCTTGCTA GTTTGGTGCT
70501 TATATACTAA AATACAATTG ATTATTAATA TTGACTTTGT GTTCAGTGAC
70551 CTGGCTAAAT TCTCTTATTA ATTATACTAG TTGTCCCATA GGTTTTCTTG
70601 GATTTTCAAT ATTTACATTC ATGTGATTTA CTAATAGTGG CAGGTTCATT
70651 TCTTCCCTTT CAATCTTGCC TTTTCTTTCC ATGCATATTG CACATGCATT
70701 GAGAACAATG TTGAATAAAA GTAGTGATAA TGGACATCTT TGTCTCTTTT
70751 TCCCGAGTTC ACAGGGAAGG TTTTCAATAT ATCAAGAGTT TATAAAATAT
70801 TTGCTGTAGG CTATTTGTAG ATATCCTTTA TCACAATAAG AAAGTTTCTT
70851 TTCTGTCCTA AGTCACTAGA AGTTTTTTTT TTTTTTAACA TGAATGAGTA
70901 CAATATTTTA TCAAATACTT TTGTTTTACT GAGGTCATTT CTATTGTGAG
70951 TGAAGCAAGT TGATTTGTAA ATATTAAAGC AATCTTGATT TCCAAAAGTA
71001 AATGCTAGTT GGTCATGTTC TATTATCCTC TTGTGTATAT TACTGGCTAC
71051 AATAAAATAT TTGTTTTTTA TATTTTTTAT ATTATTATTC ATACATTATT
71101 TATGTATGTT ATTTATTATT TATAAATATG TATTCTATTT ATATATATTC
71151 CTACATATAT TTTAGGATGT ACATAGACAA GTTTGAATGG TAACAAGAAT
71201 GAGCCAACTG AGAGGAAGAA ATTGGTAATG TAGTAAAGAG CGGGGATGAT
71251 TGCCAAGTCA GGTCCTGCAG GTGGTGAGAT GAATGTGACT CAGGGCACAG
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FIGURE 3W

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71301 GTGAATGAGC TGACCTTAGG TGGAAGTGGG GACCCTTCCT TCATGTACTA
71351 GGAGAGAAAG CAGAGTTTGA AGTCTGTATG TGTGTGAGCT GCTGGGCTTC
71401 TCAGAGGGCA GATGAAATAG TTCTTATGCC ATTGCCTGTG TTTTCCCTGT
71451 GGTATATGAG GCCATCCACT GAGAATGAAG GTGGTCAGAG TATAGGAAAT
71501 TTTGAGATGC CGAGAAGATC TGTGAAATTA GTAGAGAATT AGAATAGGAT
71551 TTTCTAAGTA TCCATTTGAG ACTTGTAGTT ATAATTAAAC AAGAATCTAT
71601 CCTGCAGATT TGTATTTTTC TCCTTAGATT GCACTTAATA GATCACCAGT
71651 TCATTTTTGT TGCTGTTTAA AAGCATATTG AGTTTAAGCA GGATTGGAGT
71701 TTAATTGGGT GAGGTATTCT CACTGTGACT AAGTTTGATG AATTGAAAAG
71751 CGTAGTTGTA GAAAGGAAAC TCAAGAAGGA AATTCTTGGG GAAACTTAAA
71801 GAATCGTATA TATGCAATGT CACTTTTTAA GACAACTAAT ATTTTTAAGA
71851 ATTTACTACT TTTGAGGTGC TGTACTAATA TATTACATGT ATAATTTCAT
71901 ATATCTTCAA CTACTAGTTC CTGTAAATAA GTATGCTGAT GATGACACGT
71951 TCCATTTCTT TCGATAGCCA CAAAAACAGG AAGTGATGAC AAAGCTGGAT
72001 TCTAACTCCT GACTCCCAAA TTCTCTAAGA CCCTCAGCAT TAACATATAT
72051 TTTATTTTAA TGTTATTATA TATGTATCAT TACTTTTACA ACTCTTAAAC
72101 CAAACATTTT AAAATTAGCT ACAACTGCAA AATCAACTTA AAAATTTCAA
72151 AGAGCCATTT AACATGATAA ATTAAAATAT TTTAGTAAAA CAAAATCACC
72201 ACTGATACTT TAATATTCTT AGGTCTGAGA AAAACCATTA TGTCGTATTA
72251 TTCCTGCGTT CCTGGTAGCG TTTCTACTGC TGGACATCAG AAATAGAGAA
72301 TAGTAGAGCC CCTGAGATAA GAGCAGAGAC AGGGGAAAAG CAAAACATTT
72351 CTGAAGAGGC AGTTGGTCTA GTTTGGCTAT AATCACTAGA CGGGTAAAGG
72401 AACATTGGGT GCATTAAAAG TAGAGAGCCT GGGATGAAGG CGTGAAGGCT
72451 GAGTAAGAAT CTCTTCACTT GGTAGTAATT CTAGTTCATC CCCCTCTGAC
72501 CTGCAATTCT GAACATGGTG TAGCTTGGTC AATAAGGAAA TAAATTGCCT
72551 TTCTGGCTGG AGAGGCAAAG GGTAGACAAT ACATTGTGCC AGCTGAACTT
72601 CCTGTCTCTC CGCTCTGGAG AAGAGCCAGT CACAATGTAT GACTCAGCAC
72651 GCCGGGCACC TCTCCCACGC CAGCCAGGCC TGCCCAGCCA CTTGCTGAAT
72701 CACAAGTGGC CATTTCCAAT CCCATCAGTG ACCCAAGCTC TCCAACTTAG
72751 ACTAGTTTCT CTGTGATCGG TCTATGATTG TCATGGAGCA CAAAAAGTAT
72801 TAACTTCTAA CATTTATTTT TCTTTCCTGG ATGCTTGATG AACTTTATAA
72851 GCAAGAGACT GATTTAATTG TTCCTCATTA TCATCTGAGC ATGCCGTCTT
72901 GGCTTGCCCT TTTATATGGA GAGCAAAATG TTGTTATTCC CCTTTGCCTG
72951 ATTACTGGCT GTATTATTCT CTGAGGTGGC CATCTCAAGA GATTCTGTAG
73001 AAAATAATAA TAGCAAAATT TCTCCCTTGA GAAGCTTCAT AAAATTAAATC
73051 TCCAGAGCCA GTATATGTAA GCCGACAGAT TATGAAATAT GATTTAATGC
73101 TCTGTCCAGA GAAAGGTCAG GGCTTCAGAA AAATCATCAT AATATCAAGA
73151 AAAACTAATC TGCAACCTGT TATATGATTT TTAAAAATCA CCCCCCATCT
73201 TTTTTACTGT GCAAACTGTA GATTTTTGTT TATTTTATTT GAGGCTATAG
73251 TTTATGTCTT GAATCACACA CATATGAGTA TTACTTTCTG TGAAGTTTTC
73301 ATGACCCCTG CAATCAAACT TGGGTCCTTC TGTTAGTTTC TATCACAGTA
73351 TCCTTCACTT TTCTTTCACA ATTCTTGCCA TATTCTATAA CTACATATTT
73401 GTTTGTTAAA TATTTGTTTA TCTTTTATAG ATGATTGGCT TCAGGAAGAG
73451 GGAAACCATG TCCTTTTTGTT CAGTCCTTTA TTCTCAGCAC CTTGCACAAC
73501 ATGAATATAC AAAAAATATT TGTAAAATGA CCATCGAATG AACAAGTGCT
73551 CATTAAGTAC CAAGCTATAT GCCAGGGGTT GCTGATGGTT AGAAATGAGC
73601 AGGGCACAAA ATTCTTTGTT CAATTAGTGA GCAATTCAGG CAAAAAGAAA
73651 ATATTAATGG TGATTATACA ATATAATGCA ATGCAGCCAT CTGCCACTAG
73701 ATTTACTGAA GTGTTTTGTT TTGTTTTTAA GAGACAGAGT CTTGCTCTGT
73751 CACCCAGACT GGAGTACAGT GGGTAAAATC ATAGCTCACT TCAGTCTCGA
73801 ACTCCTGGGC TCAAGGAATC CTCTCACCTC AACCTCTTAA GTAGCTGGGA
73851 CTACAGGTGC ATGCCACTAT ACTGGCTAAT TTAAAAACAG AAGCCAACAA
73901 ACAAAAAACA CACCTTTTTA AGACTGGGTC TCACTATGTT GCCCAGGCTG
73951 GCCTTGAACT CCTGGCCTCA AGCGATCATC CTGCCTTCCA AAGTGCTACC
74001 TTCTAGAGTA TTGGGATTAC AAGCGTGAGT CATCTGCACC AGGCCTGAAG
74051 CATTCTGTAA TGGAGAAATA CCTGGGTGCT ATGGAAGGGC AGAGGGGGAA
74101 ACACAGAGGA GTAACATCTA GTTTACGTTT GTCAAGGAGA GGCCAGGAAA
74151 GACTAACTAC AGGGGAGATA AACTCCAACC AAGAGTCTTT AAGTCTTCCA
74201 AGACTTACGT ACAAGTTTCT TATTGCTAAA ATGGAAGTTT TAATGAACAT
74251 TTATTTATTT ATTTGAGATG GGGTTTCACT CTTGTTGCCC AGGCTGGTGT
74301 GCAATGGCAC AATCTTGGCT TACTGCAACC TCTGCCCCCC AGGTTCAGGT
74351 GATTATCCTG CCTCAGCCTC CAAAGTAGCT GGAATACAGG AGCCTGCCAC
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FIGURE 3X

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74401 CATGCCCAGC TAATTTTTTT TTGTATTTGT AGTAGAGACG GGGTTTTGCC
74451 ATATTGGCCA TGCTTGTCTC AAACTCCTGA TCTCAGGTGA TCCACCCACC
74501 TCGGCCTTCC AAAGTGCTGG GATTACAGGT GTGAACCACT GCCCCCGGCC
74551 TGAACACTTA CTATAAATAT TATATGGTAG TTCTCTCAAA TTCATTCTGT
74601 TTACTGCCCA AAAGAGCTAC ATAAATTCTA AGTTGTCCAC ATTTATGAAT
74651 TTTAGATATA TGGCTGTTTA TTCTGGATAA ACACACAAAA TACACAAGAG
74701 TGGGTGCGAT CACTTATATG TGTTAAAGAA GGCATTCAAG GTGCATTTTT
74751 TCTTTGGAAA AGCTTTGTAA GGCTGCTTAT GAGACAGAGA AGTAAGTATT
74801 TTATAAATTC CAAAGCTTCT TGGTCTATTG ATGAGTTTTT CTGCTGTTAA
74851 AAACCTCTGA AAATTTGACA ACGTACTCTA GAGAGAGAAA GCGCTGAAAT
74901 AGGCACTGAC GTACTGCTGG TGGCAATTCA AAATGATATG CACCCTATGG
74951 AGATAAATTT GGCAATATCA AGCAAACATT ACATATACCT TTGCCCTTTG
75001 TTTTGACAAA TCTTTGTTTT AGCAAACCCT CTTCTATACA TCTATAATGA
75051 CATTAGACTG CCCAGAATAC AAGAAGGCAA CCACAGTGGG CCAGTACTAC
75101 TACTGGGCTA GATGTGGTGG CTCACACCTG TAACCACAAC ATTTTGGGAG
75151 GCTAAGGTAG GAAGGCTGCT TGAGGCCAGC CTGGGCAACA TAGTGAGACC
75201 TCATCTCTAC AAAAAAAAAA AAAAAAAAAA AATTAGCCAG TCATGGTGGT
75251 ACATGCCTGT AGTCTCAGCT ACTCAGGAGG CTGAGATGGA AGGACAGGTT
75301 GAGCCTTGGA AGTGGAGGCT GCGGGGAACT ATGAATATGC CACAGCACTC
75351 CAGCCTGTGC TACAGAGAGA GACTCCGTCT TAAAAAACAA AACAAAATAA
75401 CAACAACAAC AAACAAAGAT AGATGCATAG AGTTTTTCAC TGTTGCACTA
75451 TTTATATTAG CCAAAAACCG GGAAACAACC TGAATATTCA TCAAGTGGGG
75501 ACAGGTTGAG TAATCATGTG ACATACATAA ATTGCAGCAC TGCACACTTG
75551 AGAAAAGAAG TGAGAAATGT CTCTATTTCC TAGTGTGGTT TGCTCTCCAG
75601 AGTATACTGT TAAGTGAAAA AAGCACTGTG GCCTCAAATT TATCTATAGA
75651 TTCTATACAA TCCCCATCAA AATCTCAGCT GGCTTCTTTG CAGAAATTCA
75701 CAAGCTGATC TTAAAATGTG TATAGAAATC CAAGGGACTC AAAATTCAAT
75751 AAATTCAAAG ACTAGCCAAA ACAATCTTGA AAAAGAAGAG CAAAGTTGGA
75801 GGGCTCATAC TTTTCAGTTT CGAAAGTTGT TATGAAGCTA CAATAATCAA
75851 GATAGGGTGG TCCTGGCATA AGGATAAACA TGGAACAGAA TTGAGCATCT
75901 AAAAATAAAG CCTCATATTT CCAGTCAATT GACTTTTAAC CAGGGTGCCA
75951 AGAAAATTCA ATGGGGGAAG AATTTGTCTT TTCAACAACT GGTGCTGGGA
76001 CAACTGTATA TCCAAATGTA AAAGAATGAA ATTGGAACCC TACCTCACAC
76051 CATGTACAAA ATTAGCTCAA AATGGAAAAC AGAGGTAAAT ATAAGAACTT
76101 AATGTATAAA ATTCTTCGAA GAAAATACAG AAGTAGATGA TCAAGACCTT
76151 GTAATCACTA ATTGTTCCTC AGATATGACC CCAAAAGAAC AAGTACTAAA
76201 AAAAAAAGTA GACAAATTGG ACACCATCAA AATTGAAAAC TTTTATGCTT
76251 TTTATACTTC AAAGTCACTA TCAAAAAAGT GAAAAGTCAC CCCAGAGAAT
76301 GGGGAGAAAA TATTTGCAAA TCATATATCT ACTAAAGGAT GTGCATTTAC
76351 AATATACAAA GGGGCCAGGC GCTGTGGCTC ATGCCTGTAA TCCCAGCAAA
76401 TCGGGAGGCC AAGGTGGGTG GATCACCTGA GGTCAGGAGT TCAAGACCAG
76451 CCTGATCAAC ATGGTGAAAC CCTGTCTCTA CTAAAAATAT AAAAATTAGC
76501 TGGGTGTGGT GTCAGGTACC TGTATCCCCA GCTACTTGGG AGGCTGAGGC
76551 AGGAGAATCA CTTGAACCTG GGAGGTAGAG GTTGCAGGGC GTGGAGATTG
76601 TGCCATTGCA CTCCAGCCTG GGCAACAAGA GCGAAACTCC ATATCAAAAA
76651 AAACAAAAAA AAACAAAAAA AAACAAAAAA AAAAAAAGAA CAAAGATTTC
76701 TTCCAAGTCA ATAATAAAAA CAGAAAATGC AATTTAAAAA TGGATAAAGA
76751 ATCTGAGTAG TTTTACATTA AAAGATAAAT AAATGGTCAG TGAGCACTTC
76801 AAAAGATCCT GAGCATTACT AAACATTAGA GAAATGCAAA TCAAAATCAC
76851 AATGAGATGT CATTTCATAC CTATTGCTTT CTTTTTCTTT TTTTTTTTTT
76901 TTGAGACAGA ATCTTGCTCT ATCTTCCAGG CTGGAGTGCA GTGTGTGTGA
76951 TCATGAAAAT GGCTCACTGC AGCCTCAACA TCCTGGGCTC AAGTCATCCT
77001 CCTGCCTCAG CCTCTTGAGT AGCTGGGACT GCAGGCATGT GCCACCGCAC
77051 CAGACAATTT TTTTTTTCTT TTGTAGACAC AGTGTCTCAC TATGTTGCCC
77101 AGGCTGGTCT GAAACTCCTG GGTTGAAGCA ATCTTTCTGC CTCAGCCCCC
77151 CAAAGTGCTG TAAGTATAGG TGTGAGCCAC CACACTGGGC CAGTACTATT
77201 CTTTAAAAAA TGGGAAATAA CAAGTGTTGG AGAGGATGTA GAGAAACTGG
77251 AGCCTTTGTA CATTGATAGT GGGAATGTAA TGTGGTACAG CCACTGAAGA
77301 AAACAGTTGG ACAGTTCTTC AAAAAGTTAA ACATAGAGTT TCCATTTGAT
77351 CCAACAATTC CGTTACTCAA TATTTACTCA AAATAATTGA AAGCAGGGAC
77401 TCAAATAGAT ACTTGCACAC CAGTGTTCAC AGCAGCATTA TTCATAATAG
77451 TCAAAAGGTA GAAATAACCC GAATGTCCAT CAACAGATGA ATGGATAAAC
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FIGURE 3Y

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77501 ACCACATAGT ATGTGCCTAT GATGGAATAT TACTCAGCCT TATAAAGGAG
77551 TAAAATTCTG ATATACACTA CAACATGGAT GAACCTTGAA ATCTTATAAT
77601 AAATGAAATA ATCCAGACAC AAAAGGACCA ATATTATATG ATTCCACTTA
77651 GATGAGATGC CTAGAACAGA CAAATTCATA GAAACAGAAA ATAAAATAGA
77701 GGTTACCAGG AGTTGGAGAG GAGGAATAAG GAGTTATTAT TAAATGGGTA
77751 TAGAGTTTCT GTTAGCAATG ATGAAAATGT TCTAAAAATG GACAGTGGTG
77801 ATGGTTGTAG AACATTCTGA ACGTACATAG TGCCACTGAA TTGTACTTAA
77851 AGTGGTTAAA ATGATAAATT ATATGATATG TATATTTTAC CACAATAGAA
77901 AAAAATACAA GAAGTTACCA GTGGGGAAAA GGAGGGATTA CAGAAGACAG
77951 GGATAACAGC ACGACTTTTC TCAGTATACC TTGTTTTTCG TATTTGACTT
78001 TGAAAATATG TACATACTTT ATATAACTAG AAAACAAAAT TAAATCTTAA
78051 AACAATCCCA AAAATGGAAT GTAAAAAAAA TGAAACCAAT TAATCTAAGT
78101 ATATATCCAG TTTGTGGCAT AACCACACAA AAATGAACTA TTCCAAGTGA
78151 CTTTTGAACA GAAAATTACT ATATACCATC AGTAGAATAT ATCCTAATAA
78201 CAAGAAAGAA CAGCAAAAAT ATCTTAAAGT GTTTTCAGTA ATGGCATTGT
78251 TGGGGGTAAT GTTGATACTG TTATTTTGAA AGTGTTGAGT GTATACAGTG
78301 GGATAGAACC AACAAGTATT TATAATGATA TCATTGAGAA CCAAGATTTT
78351 CATTGAGGGA GAAGACTGAT GAAGTTAAGA ATTTCTGTAA TCTTGAATGT
78401 AAACTGAAAG CATTATTATG AAATGTGTGA TGTGTTTATC TTAGTTTACC
78451 TTTGAATATG TGTATATTTA TAACTATACA TCTATAGCAG CAGACACTTC
78501 TGTCACCCAG ATTGTCTGAA ACAGGAAATA TACAAGATAG CCAGCAATAT
78551 GTTTTCATAT TCTACAGTTA CAAAGCTGTC AAAACTTACT AGGGTTATGT
78601 CAAACAAAAC ATGATCTAAC ATGACTATGT TCCTACTGGC TGAAGAATGA
78651 ACATTATGAA CTGAACATCA ATAAGAATAA TGACATCAAA CCCAGGAGTT
78701 CATTATAATA TATTTTTAAG TATATTGATT GCTTTTTGGAG GGTTCTAGGA
78751 AACAAACAAA TCATTTTGAA AAGTGGTAAA TAAAGGAAAG ACTTCAGTTC
78801 AAGACCAGTC TGAGCAACAT AGTAAGACCC CATCTCTACA AAAAATTAAA
78851 ATATCAGCTG AGCATTGTGG TGTACATCTT TAGTCCTAGC CACTTGAAGG
78901 CTGAGGCTGG AGGATTGCCT GAGCCCAGGA GTTCAAGGCT GCAGTGAACT
78951 ATGATGGCAC CACTGTGGTC CAGCCAGGGT TAAATAGCAA GACCCTGTTT
79001 CTGGCGAAAA AAAAAAAAAA AAAAAAAAGG AAGACTTAAA CATACCTTTC
79051 CTATATGAAC TGTGCCTCGG AGTAACTAAA TAATTGATTA AAGCAAGTTT
79101 CTCTGTATAA AAGTACTCCA GCTAAAACAT TAAGGAGAAA TGATAGAATT
79151 CAAATATCAC AACCCCTAAG GAATTTTTGC ATCAAGACAA CAATAATTAA
79201 TGACTGATAA CACCACACAC AGAATACAGA CTTATTAATT GTATAACTCC
79251 TGATCAAGTG CATACCACTA TCTGTGAAAT AGTTTTGCCA AAAAAAAAAA
79301 AAAAAATCTA ACCTAAACTT GAACAAGCCT CTAGATCTAA CCACCAATTT
79351 TTACAAACTA CAAAGAATTG TGGAATGTAT AGATTGACGT GACATGAAGG
79401 CAATCGGCAA AGTCCAGACT GTGAAAATAC TACAGCAAAC ATTTAGGGTC
79451 TTTTTTTCTT TTTCTTTCTT TTTTTTTTTT TTTTTTTTTT TTTTGAGAGA
79501 GTCTCCCTCT GTTTCCCAGG CTAGAGTGCA GTGGTGTGAT CTCGGCTCAC
79551 TGCAACCTCC GCCGCCCAGG TTCAAGTGAT TCTCCTACCT CAGCCTCCTG
79601 AGTAGCTGAG ATTATAGGTG CGCGCCACCA TGCCCAGCTA ATTTTTGTAT
79651 TTTTAGTAGA GACGGGTTTC ACCATGTTGG TAAGCCTGGT CTCAAACTCC
79701 TGACCTCGTG ATCCACCCGC TTCAGCCTCC CAAAGTGCTG GGATTGCAGG
79751 CGTGAGCCAC TGCACCCAGC CCACCCTTGG TTTTTTTTCAA CAAAAAATTA
79801 CTAGAAATAA AAGAATAATA GTTGGTCAAG GAAGCTGTAG AATAAGAAAG
79851 ACTGCCACAT ACATCAATGG CAGTGGGCGG GCTTTGTTTG AATCCAACTC
79901 TAGCATGCAA ACATTTGATA AAAATTTCTT TATTTAAAAA GAAAAGTTTA
79951 CAAAACAATC AGAAAAAATA AAAAAGATTG AGGATCTCAG GACAACTACT
80001 AGCCTAGATA ATTTATAAAG ATTAGATAAC TGACTCATTT TTATTAGTTT
80051 CTTTCCTAAT AAGGCAATAT GTATTAGATA TATCAGAGTA GAAGGAAATA
80101 TTTTTCTTAC ATCTATTTGG CTTTTTAAAT ATAAACATAT ATAAGTAAAA
80151 ACCAAAATGA TTTATAATCC CACCATTTAT GTAACTATCT TATTTTCAAA
80201 AAAAATTATG CAAATACTAG CATTTGTGTG CTTTTTTTCC TTTTGTGTTT
80251 GTGTGTTTAT ATCCTTTTTA AATATATCCT TTTTATGTAC CTAAGCAGCT
80301 GTATACTATA CTGCATACTA TAGTGTGAAC TTTGTTCTTT TCCTTCGTCT
80351 TTACAACATA TTGTGGAAAA CGTTCCATAT CAGAATATAG ATATGCCTTT
80401 TTGTAGCCAT TGAAATGCAA AGAAAAAAAG AATATAGATC TGTCTCATTT
80451 TTTAAAAATG CTGTATAATC TGTAGCACGA ATTTACTATA ATTTATTCGC
80501 ATGCTCCCTT ATCGATGGGC ATGTAAATTG TGTTAATTTT ATATGATATA
80551 ATGAGTATCC TTATATGTAT ATCTTGGCAC AGTTTTTCGA GTGTATCCAT
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FIGURE 3Z

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80601 AAAGTTTCTT GCAATGAAAT TATAGGGCAA CAAGGGTGTG GTGGCTCTTG
80651 TCTGTAATTT CAACACTTTG AGAGGCTACG GCAGGAGGAT TACTTGAGGC
80701 CAGGAGTTTG AGACCAGCGT GGACAACATA GTGAGCCCTC ACCTCTACTA
80751 AAAATTAAAA AAAAAAAAAA GAAAAAGTTT GGTATGGTGA TATGTACCTG
80801 TAGTCCCAGA TACCCAGGAG GCTGAGGTGG GAGGATCATT TGAACCTGGG
80851 ATGTCAAGGC TACAGTGAGC TATGACTGTG CCACTGCACT GCAGCCTGGA
80901 TGACACAGTG AGACCCTGTC TCAAAAAAAA AAAAAAAAAT TACAGGCCAA
80951 ATCCATATGC TTTTAAAGGA TATTTTTGAA TTGTTCTCAA AAAGAGGCTT
81001 CACCAAATTA CCATCCAGGG TATACAAGAT ACCCATTTCT CCATGTCCTT
81051 ACCAACAGTG GCTCTCATCA AGCCTTGGTG GAAATGCTCT CATACTGATA
81101 CTTTAACGAC TAAAAGTCAT GACATATCTG CTTAGGTTGT AAATTGCCTC
81151 CCTCTAAACT TATACAGAGA GAATTTAGAG TGTTGTCTCA GCTTGGTTCC
81201 AGTGTTATCC AAGCCATTAA CCTTTGTTTT GCCTTAGATT GTCACATTGT
81251 GGTATTTCAG TTAAAAAACA AAAACACAAC TGGTACTTTT TTTTTTTTTT
81301 TTTTTTTGAG ACGGAGTCTC GCTGTGTCGC CCAGGCTGGA GTGCAGTGGC
81351 GTGATCTTGG CTCACTGCAA GCTCCGCCTC CTGGGTTCAA GCCATTCTCC
81401 TGCCTCAGCC TCCCGAGTAG CTGGACCTAC GGGTGCATGC CACCACCCCC
81451 GGCTAATTTT TTGTATTTTT AGTAGAGACA GGGTTTCACC ATGTTAGCCA
81501 GGATGGTCTC GGTCTCCTGA CCTCGTGATC CGCCCGCCTC GGCCTCCCAA
81551 AGTGCTGGGA TTACAGGCAT GAGCCACTGT GCCTGGCCAC AATGGGGTAT
81601 TGTTTTTATA GACTGTTGAA ATCTGCCTTT GGAAACCATG GGTTTGCTGT
81651 GTTGTTATGG TGAATGAATT AGGTGCACAA TACTAGTTTT TAAAAAATGA
81701 ACTTCACACT AGGTACACCT TGAAAAATTA TTCCAGAGCT ATAAGAAGAG
81751 CTATAAGAAG AAAAATATGA TGGGTCATTG CTCCAAAGAA AGGTTTTAAA
81801 ATGTAAATTT GTACTTAATG AATAGGACAG TGTACCCTAA CCTCCTCCTT
81851 GCTATTCTTC AGGGATCTCT TCTAACAAGG GCTAATGCTT CACCTAAGCT
81901 GTGAAAAGCC TGCTGTGAGC ACTCCCTGTT CAGGGTCAGA AAAACACAAT
81951 GAACTGTTCT ATCATTTTAG GTTCTAGGAC AATGTTCTCT TGCTTTTCCT
82001 TGCTCAGAAT GGACCCTTGC TGGGGTAGCA TCAGAATGAG GATCTGGTGC
82051 AACAGTTCTG CAATAGGAAG TAGGTTCCCC TACTATCATG GTTTTCAAGC
82101 TTTTTTGACT GCAGCCCATA ACGAGAAATA ATGTTTTTCA TCATAACCCA
82151 GTAGATATAC TCACAGAGAC ACAGTATATT CATAAAAAAA ATCATAACGT
82201 TTAACCTTAT GTTAATAGCA TTTATCCTAT GTTATTCAAT CTATTTTATT
82251 TCTTTTTAAA AAATGCTCAT CACAGTTAAC TAAACTGATT TCACAACTCC
82301 TTAAAGGAAT TTGACTCACA ATTTGAAAAA CACTGCATTG TAGAATATTT
82351 TAGAGTCTCT TCCCAACCCT CAGAGTCAGA TTTATTTCAA GATGGCCCCT
82401 GTAAGACAGC TTCAAGCTTG TGAGTGACTT TCTTTTTTCT TTTTACTTCT
82451 TTACCATTTA CCATGACTCC CAAATAAGTG ACTCTTTTGG CTTATTTGGT
82501 AACCATGCTA ATTTCTACAC ATAGAACCTA GAGCATTTAC ATAAGACCCA
82551 CCCAAAGCTT GTGTTTTAAC CTTGCTTCTC TCCTTTCTTT CTTTGATTCA
82601 TTGATTATGT TTTCTATTGC TATCTGTTCA ATCTGTGTTT CAGGCAGTGT
82651 ACAGGTACTG AGGCAACAAT GGTGAGTAAA AGCAAGCATG CATCCTGAGA
82701 TATACTGGGA ATGAAAGAAG CTAATCCAAA AGCATACAGG AAAAATATTT
82751 CAAACTTTGA TAAATTCTGT GTAAGCATAT GGCATTGCAC GTAACAGGGG
82801 AACCGCATTT AATATGGAGT GTTGGAAAAG GCTTCTGTGA GAAGTGACAC
82851 TTGAGCTAAG ACTAGAAAAG TGAAAAGAAT ATAACCAGGT ACTGGACAGC
82901 ATCATGAGTG CAGGCACAGG TGACATCGTA TCACAAGCTT CTAAGGCTGA
82951 AGGGGGCGTG AATTGCTAGC TGGAGAGTGG AAGGAAAAGA TCTTCAAGAT
83001 AAAGCTGGAA AAATAAACAG GGCCAGGCCT CATAGGTTTC TGTAGACCAT
83051 GGAAAGAGGT GAAGGTTATT TTGAGCCTGG ATGACATGAT AAAACTCACA
83101 TTGTAAAAAT ATAACTGCAA GGTAGAGAAT GGATTGAAGA GGTCCAAGAT
83151 TACGCAGACA GAGCTATGAA CAGCCTATTG CAATGGTCTG GGTCAAGCAT
83201 GATGGAGTAG GGTTGGAATA GGGTGGTGAA CTTTTATTAG TTATCTTCCT
83251 TACTGAGCAC ACTTTGCAAT GAATTTCAAA TGCACTGGGA CCAGACTTGT
83301 TAATTTTGGA GCTGTCGACT AACAAATAAG TAAGCCATGA TAACCCACCA
83351 AAGAAAGTTG CAGAAATGCA AGAGCAAGGC TGTGATGAAT GGTTGAGGTA
83401 CAAGGAAGCT CTTACTCACT CATTTTAAAA AATCAGATGA TATGAAGTTG
83451 AATATTCAAG ATATTGCCCA ATTGTGTTAT GTTCACATAT TTTACTGGGC
83501 ATAGTTCTGG ATAATAAAAT ATTTATCTTC TCTCCCTCTG AGAATTAAAA
83551 ATCTGAGATG GAGGCCTCTG ATGTGCCAAA GGAGAAAGAT GATTTTTAAG
83601 AGCCAAACGT GCCTCCATGA TTAAATACAT TTATATTTCT ACTGGCCAAG
83651 GAAAGCATGT TGCCTCTTGC CTGGGCCTCT TCTGTCTTTG ATTAATAATC
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FIGURE 3AA

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83701 CCCTGCACAT TCGAACACTG TTATTAACTT GCCACATTGG CACCTTTATC
83751 ACTTTGTTCT TTGAATAAAA AGAGCTTAAC CCAAGTCCCA GTAAAAATGT
83801 TCATTCAGGC TGAATTTAAG AAATATATTC TGCTCCCTTG GAGTTAAATG
83851 GAATAATAGG AGAAGAGTCC ACTTGACTGT TACCAGGTTT CTGAACTACA
83901 CCTGGCAGCC TAACATAGTC AACAGCAGGG AGTGAATCAC ATCTGCTCTG
83951 TATGCTAACC CGGTCTGAGT AGGTGGTTTG CATTGGCATC TAATTATTTT
84001 TATGGTTAGT ACTCTCTTCT CCTGACTTTT GGTACCAAAC CCTCACACAC
84051 CTCATTATCC CTATTGCATC TGCCACTCAT CCTAAAAGGC CTTGCTTACA
84101 TCCCACAATC AATCATTCTT TCTCTTACCT TAGCGGAGAA CAGCCTGAGG
84151 TGCAGCAGGT CCCAGATATG ATTACAGTTT CACCAGTTCA ATATTGTTTA
84201 CTGAATGGCC TGTAAAACAC AGTGAATATA ATTTGTGTTG CTGCAGTTGG
84251 AAGGCTTACA TACCACATTG CCTAGAACCA AAGACCTTTC CTCATGCCCA
84301 ATACACCAAT GGCAGAGATG ACCAGCCAGT CACTGCATCG AGATGAAGAA
84351 TAGTATCTCC CAAAAGGCAA TACCAAGCAT ATGTTTCTCA GGCTTTTACA
84401 AAACACTTTT TAAGTTTCTG TCTAAACTCC TCTAAGAGCT AAATTTTTCC
84451 AAGACGTATT CTGTGTAAAT CAGTCTTCAG TGATAAACAA AATTTTATTT
84501 ATTGAACTAT CAGGTGCTAT TAATGCTAAT TAGAATGTTA CCACCTCAGA
84551 TTAATGCTTC GTTGAATTTC TTTTTTTTCT GGTGTTTGTA AGTATTCCTT
84601 TTCTCCTTCA GCACAATGAT AATTATAAAG AAGAAAATGT ACTAAGTGCA
84651 TTTCTCCCAT CATTTGATAT TTTACATTTA TTTCCTCAGC AAATAATTTG
84701 TCACAAGGAA GTAATGTGCA TCCCTGGGCA CTGCTTGCAG GCACTTAATT
84751 CTTGATTCAA ATGAAACTTT AAAATGTTTT ATCCATGATG TTATGTCTAA
84801 AGAAACATGT CAAAGAAACA TGTCAGAGAA CTTGACTTTG AATAGAAATC
84851 ATGGCTGTGC TTTGAGGGAA ACAAAATAAA TCACAGAGGT AGGAATGCAT
84901 AGTTACAAGC TACTGTTTGT ACACAGCAGA GACCAATTCT ACTCTCTGTT
84951 CTCATTTCCT CTTCTAATTC CTCATCCCTA CACTCCTTCC TGTGTGAAGC
85001 CCATGTCTGA TCCTGCCTAA TTCAGTGACT GGGGGTCACT GCAGATGCGT
85051 GCACAGGGTC CTGTTATGGG ATCCGGATTC TGCCGCCTTC TCCAGACACA
85101 AGTTTCCCCT CATACCTGTT GTTCCAGCAA ATCCAAGCTA TTCTCCTTTC
85151 CCCACTTGCA CTAGGTTCTT TCCCTAGTCT GTGCTTGCAT GCATCCTATT
85201 TTTCTCTGGT ATTTTTCAAA TTTTACTTTG GCACCTGGAG AACGTTTTGG
85251 CACCACCATT TGTCAGGTGT TTAACTTTGT GCATTTCCTC GTGTGAATGG
85301 GAGCGTAGGT CCAGCATCGT GAGGAAGGAC TGGGGTCACA CTCACAGAGT
85351 GTGTCAGAGC CCACAAAGTC ACTCAGTAGA AACATCAGGA GATGTTAGCG
85401 TTATTTTTCA GTTATTACTA TGATCACCAT TCCTCAAAAT TGAGCTCTGG
85451 TTTTACCTCT CCTGACAAGC TTTCCTTTAC TTCCCCATCC CAAAGACAGA
85501 GTGAATTACT TCCTTGTACT GTGTGCTTAG TTCTTCATTG CCCTTCTTAT
85551 GTGTTTTCCT TATCATTAAT GTGGGACATG ATCTGTTATA ATGTTGCTGG
85601 GCAATGATGT TGTTAGTATA GAAAAATGGG CATGAGGATA GTTCAAGGAG
85651 TTCCCATAAC TCATATTTTA TGGGCCTTCT GCAATATATG GTTAGGATAC
85701 AACCATTAGC AATAAATGGA TAACTTGGGT TCTCTTCATT TTCTGTGTTT
85751 TATTGCTACA TGAATAAACA GTTATTGAGT GCTTACTGTA TGTCAAGCAT
85801 GACAATAAGT ATTATAATTA CCCTGTTTAT TCATCAGTAT GATCAAATGT
85851 GGTTATTATT CCCATGTGAC CCATGAGGAA ACTAAAGGCC TAAGGTGATA
85901 GAGCTAGTGA TAGACCACCT ACTCCCAAAG TCTGAGCTCT TAGCTCAAGA
85951 ACACTCTGCT CTGATCTGTA GGGTCTCATT TGTCTCTGAG ACTCTTTAAT
86001 GTGTAAATAT ATTTGATAAG TTTTCTCTTC TAATGTAATT CCAGGTATTC
86051 CTTCCAAGAT GAGGAAGACA TGTTCATGGT GGTGGACCTC CTGCTGGGTG
86101 GAGACCTGCG TTATCACCTG CAACAGAACG TCCACTTCAA GGAAGAAACA
86151 GTGAAGCTCT TCATCTGTGA GCTGGTCATG GCCCTGGACT ACCTGCAGAA
86201 CCAGCGCATC ATTCACAGGT CAGTCAAGTC CAAGGAGATG GCCATGAACG
86251 TAACGCAAGG AGAGAATCCA CAACTGGCTA CCTTCAATAA ATTCTTATTG
86301 AACATGACAT TTAATCCCCG TTTAATTCTT GAAACAGTAC CCTGAGGTAG
86351 GTTGATTGTC TTCATTTTGC AGATTTTGTA AAAGACTGAA CACATAGAGC
86401 TTAATTTGCC AAAGGTCACA GTAAACAACA AGATCACAAT CAATGAATTT
86451 TGGTACTATT TTATAACTAA GCTTAGACAA AAAGGAGAAA AGGTGACATA
86501 TAGAAACCTA ATAAATATTA AGTAAATAAT TAAATGGAGG TAGCACATGG
86551 AGGGAAAGAA ATAGAATGAA AAGAAAGAAA GTTCTTTGGG AAAAAAGCTT
86601 GAGTCTTTCT AATATTTGCT GTCCTGCAGT CTATATTAAA TTAATCCCTA
86651 ATGTATGTAC TGCAAATGGA GGTAGAAAAA GCAATAGCAA TGTCTTCTGC
86701 ATTTAGAGCA TTAGTAGTAA ATAAAGACAT ACAAATAACA TAAGAAACCA
86751 TAAAGCTATA GAGATAATAC AGAGAAAAGG ATAATACTTT ATAGTAAAGA
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FIGURE 3BB

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86801 AATTTGTAGT TTCAATGATG ATTTTATATA TAGTATCTCA TTTGATCTCT
86851 GAAATAACCT GAGATAAATG ATCAGAGCAG ATATAATTAG ACTAGAATTA
86901 CATATGAAAA AATCATGGCT TGTATACATT AAATTATCAC CCAGTTTACT
86951 TATATGAATT GTAAACATAT CAAACATCAA AACATCTACT AATCAACATC
87001 AAAACAACTA GTGTTTACTG GTTGATGACT TACTATGTGC CAGGCACTCC
87051 TAGGTACTTT ATGTACATTA GTTTATTAAA TCCTCAAAAC TCAGCAAAGA
87101 TTCCACATTT CATTATAATA TTCCCATTAC ACAGATAAAG AAACTGTCTC
87151 AAAGGTTTGC CAAGGACAAA CAGCTAACAA ATAGCGTAGC CAGGATTTAA
87201 ACCTAGATCT CTCTGACCTC AAAGTCAGAA TTCTATGATA CCAATTCACA
87251 TTACTTACAC ATATGAAATA TATGCATTAA TTGATTATAC ATCATTAAAT
87301 GAAAAATCAG TACATGTGAC TCTGCTGCTG TCATCTCTAA TCCTTGAAGA
87351 ATTTGCTGAG ATTTTAAGTA CAATTATGTC TCAATTAGTA AAAAGTTGGC
87401 TAGATAAAAT ATTTGACCAC CACCAGTTGA CATTGACCTG TAATTTATTT
87451 TTTAAACCTT TATATATATA TATATATTTA GAGAGATGGG GTTTCACCAT
87501 GTTGCCCAGT CTGGTCTCCA ACTTTTGGCC TCAAGTTGTC CTCCTGCCTC
87551 AGCCTCCCAA AATGCTGGGA TTACAGGAGT GAGCCACTGT ACTCAGCCTA
87601 TAATTTATCT TGATGAGTAC AGAGCCTATA GATGAAGGTG AAGCATCAGA
87651 ATTTATAGAT TCTCTGTGCA GGTACCACAG GCCAGTTCTT TTATTTATTT
87701 TTATTTTTTT GGGCCTTGGC CCTCTACATT TAGTTTTTAT TTAATGTTCC
87751 TTCTTTGGAA GGGCCTGCTT GTATTGGAAG TGTGCTCTTC AGGCACCAGA
87801 TAAATGAAAG CAGACCAGTT AATTACGTAG GATCTCAGAA GTGAATTTGC
87851 ACACCTGGTG TTTTTTTCAA TAACTAGAAA TCCTGTTCTC AAGCACTCAT
87901 CTTCCCATAC TGGTTTTCTG GTCCCTCATA GCTCTTTCTG AAGAGAGACT
87951 GTTCATACTT GTTAGTCTAT GGAGTCCCTC TCAAAACTTT CCTGCTCGTT
88001 CATTCTCCCA AAAATTGCCA ACCACAGCCT ATCTTGGTTG TGACATCACA
88051 GATATCAGAA AGAAGGCAGT GACCTTGAGA AACCAGCATG GCCTCAGAGC
88101 CTTTTCACTC TCTCTCCTTT TCCTGTTTGA AATTGGGTTC TGTCCCTTCT
88151 TTCTTTAGGC TTCATGTTCT TGGTCATCAA AAGACCAATT CTCTGAGCAT
88201 TTTCTCCATG TACTTAGAAC TGTGTTCCAA GAGGAATTCA GGAGGGAAAA
88251 ACAACAACAA AAATATTGAT ACAATTTTTC CCCAAGGAGC TTACTAACAC
88301 CCAATACTGT TTTTCTGTTC TTTCCCTCTC TTTTTTTCTC ACCGTTATCA
88351 TCATTTTGCC ACTTAAATCA TAAACCAAGG ATTAACTTTC TGGTTTTTTG
88401 CCCTTCAATC ACATCCACAG TTATTACTTA GTGCCCGTTC TCAGAAGGGC
88451 CTTTTTGTAC TGAAATGTCT CCTCACCATG GTAAAGGTAT GGAAGGCAAA
88501 CAGGATGACA TTTTGAGTGC AGTGTTAAAT TGAGGTGACA TCCTTCTGGT
88551 GTCAAAAACT ATTCAGGTGC ATTTCTGTAA CCTCTATGCA CCTCTCCCCC
88601 CACCTCCCAG GTGTTATATT TTACAGGCTG TCATACCCTT TTGTACCTCT
88651 CCTGAGGAGT TGTGACATTT GGTGTATAAT TAATTCATTT GTCTCCTTTA
88701 TAAAATTGTG AACTCTGCAT GTTTTGCTTT TCATTGTATA ACCAGTATGT
88751 GAAAAAAATA TGAGCCACAT GAATGAATGA TTGACCAGAA GTTCAGGCTT
88801 ACAAGTAGGA AATATTCAAA TATAGGACAT TAAATCCAAA GGCCTCAGAC
88851 CTACTTGTAC CTTGGTCTTT ACATTAATCA TGTTATTTAT CATCCAAACC
88901 AGGATACTCT GAGAGCTAAA GAGGATGCTA TTAATATTAA TAGCACTGGG
88951 AAGAGTCAAA AGCCATAAAT AATCTAGGCA ATTCAGGACC TATGTCAACA
89001 TCATTAAGGC TTTTCAAGGC AGTGTTTTTT GGTTTTTTAT TTTTTGTAGA
89051 GACAGGGTCT CCCTATGTTG CCTAGGCTGG CCTTGAACTC CTGGGCTCAA
89101 GCAATCCTCC TGCCTCAGCC TCCCAAAACT CTGGGATTAC AGGTGTGAGT
89151 CACCATGCCC AGCTTCAAAT AGACATTTTA ATTCTGACAG TGTTCTGATA
89201 ACCAGGATTT TCTGCTCTCA GAATACCAGA TATCAATTTG AAATGGTGTC
89251 AAATAGCTTT TTAAAAAGTG TACATGGTAA AAGAAGCAGT GATCCCTTTG
89301 TTTAAGGAAT TTAAATGATA ATAACTTTGT CAATCTGAGA CTAAGAACTC
89351 CTGGGCCAGA GAGTGCAAAA AGCAATACAG AAGAGATACA GGCTTCTGAA
89401 TACTGTAATT CTTTTTTAAA CCTCCTTCTT CAAAAGAATC AGCCCGATTC
89451 ATGTTGTACT TGAATTCAAG ATAACAAAAC ACCTTTTAGT TACTTAGAAA
89501 GATTAGATTG TAAAATATGT GCTGAGTTCC TAGAAATTAA AAGTGAGAAT
89551 GAAAAAAAGA ATCAATGAAA GTACAGTAGA TCTCCCGGAC AAGGAGAGAC
89601 CATCTGCATA AAACTGAAGA TATAAAATAT GTGACTTCCT ACTTTTAGAT
89651 TAAAATCTAC ATTTTGCCTT TGGACATGGT AGAAGATTCA AAATTACCCG
89701 TAAACAGTCA GCACTACGTG GAAGTAGGAG CAGCAGTAGG CTGCTGTTTG
89751 CTTAGGGTTT CCTGGGTACC AGGCTGCCTG CTAAGCACTT GTGAGTTATT
89801 TCACTCAGTC TTCCCATAGC TCCAGGAGGT TTATGGCACT TTGTCCCCAT
89851 TTCACCTTCG ATGAAACTCT GGTTCTGAAA AATTACTTGC CCAAGTTTGC
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FIGURE 3CC

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89901 ATGGCTATTA AGTAGGGAAA GCATCATGTT TAGGAAATGC AGAGCTCTTC
89951 ACCACTCTCC AGCCTGCAGA TGCTCAGCAT GGCTGCAGCT CTGAGGGGAG
90001 CGCGGGACAC CTATGCATGG CCACCTGCCT CAGGCACCCA CAGACGAAAG
90051 TGGTACATGT GGAACGGACA GACAGAGAAC AGCCTAAAAT TGGAAGCTAA
90101 ATTGTGTGAG AAAGACAAGT ACTTCAGAGA AGATAGTGTG GAGTCGCAAA
90151 ATAAGTTTCA TGAGAGCTCA TACAGAAAAC AGCCTAAAAC TAGAAGCTAA
90201 ATTGCGTAAG AAAGACAAGT ACTTCAGAGA AGTTGGTTGG GAGTAAGAAA
90251 GCAAGTCTCA TGAGAGCTCT GAGGGGTGTA AATGGGACTT TTAACAGCCA
90301 AAGCACACAG CAAGTCTAGC CTAGCAAGAG GAGCTCAATG GATGGAAGTC
90351 CTCACTTGTT TCCCTGTGTT AACATAGAAG GGGGTCTTTT TAAAATTTTG
90401 TTTTCACTTC AGCTTTTCTG CCAGAAATGT CTAGTGTAGT GATGTTTTAA
90451 AAAAAACCTA AGTATCTGTT TCCGCCACAA ATCCCCATTA AGACATAAAT
90501 GGAGTTTTAT TTTGTGGATG TTTAAAAATC CATGGACTTG AACTTTTGGT
90551 AGTTTCCCAA ATATGTAGAA TATTCAGCTA GTTTTCTTCA ATTTCAGAAT
90601 CTTTCTTTTC TATCATTGTT AAAGACACAG GGTTGCATAA TAACCATTAA
90651 GTTTGAATTG TGCAATTAGA CAACTTTCTT ATTAGTCAAG AAGTCAAACT
90701 TTTTGTGTGA GTACAGCTTG AAAATCAGCT TTAGTTTCCA AAGAATGGCC
90751 AGTTTGAAGT ATAATATTCT CTTTTGCTTA CTTGAAATCT GCAAATAAAT
90801 GCTTTAAATT AGGGACAAAG TGATTATTTG CTTTTATTTA AAAAATAAGG
90851 GAAACAAAAC TCATTACAAT CTCTTCTACA GGGTTAGTAC TATTCTATTT
90901 GTTGATTGCC TCAGCCTCTC CAATGAACAA TCTGGTGGAA AGTAATTATT
90951 TAATATTATA ATCCAAAGAC AAATTTCTGT TTACTCCCTT GTCAGATCTT
91001 AAAGTAGACT CAATTATGAA TTTAAGCTAA TGAGATGGAT TGTATGGGAC
91051 AATTAAATAG TAAGTCATTT TGGGTCAAAA TACCATTTGA GAGGATGGTT
91101 GATTGTTTTT TCCCTCTGAG AATTACCCCC CACTATAACG AGGTTATAAC
91151 TCACTGTTTG CTAAATTTTT ATAGGAATGA GATAAAAAAT CTGATTAGAG
91201 TAATTTGTGC AAGTAATTAC AGTACAACAG AGAGAGTTGC AAAAATTTCA
91251 TTTCCCATTG AGTACCGAAA TGTTGAAGAG AAATAAAAGA AGATTTATGG
91301 CTGTGTAGAA AAACACAGGA TGGTATTTTT ATTTATCACC TTTGCCTTCT
91351 TTGCTGTTCT CATTGGAACC AATAACTGAT TCCAGATTCA TCTTAGGGAC
91401 TGTATAAGAT GCAGATAGAA ATTATTTCTC ACACATGACC TCTTGGGCTG
91451 GAGTAGCTGC TTATGAGATG TTCCTATCAT TCTTCTAGAA ATCAGTACCT
91501 TGACAGTGAA GAAAAAAATC TTAGGAATAA TGCTTCTAGT CCAAATATTT
91551 ATTCAAAAAT TATTTACTGG GTACCTATTT GCCAGTGTTC TGAATGCCAG
91601 GCTCCCATGG GGAAGAAGAC AATCCCCCTG TCATAAGAAG TTGTTAATAT
91651 TATAGTGTGA AAAATAGTCA AGTAAACACT TCAACATTAA TATCAAAAGG
91701 CTTTTAAATG TTGTGGCATG TGCCATAAAG AATGAAAGCT GTTATGTGCA
91751 TATCCTGAGC GATGCATGTG TGCCTGCATG CACGTGCACG CACGCACACA
91801 CACACACAAT ATGCTTAGTT GCGTCTTCCC AATGCTCATG GTTATACCTC
91851 TAATTGTAGC CTCTGGACCA TGATATTCTA TATAAAAAGC TGTCTCCCCT
91901 CTCCAATCTT AAGCCCTCAT AAGTGGATAC TACACCTCAC TTATGTTTTA
91951 ATCTCCAGCA ACTTGCACTG GATCTAAACT AGAGTGCTTG CTGGATAATT
92001 CAATGACTGA ACAAATGAAT GAGGACAGTA TGTATATGTA ACCATTGGGT
92051 GAGTGCAGAA GGTAAAAGTT GCTGTGGAGG ATGTCGTCTT CAGCAAATTC
92101 TCAAATTTAT TCCACACATT CCTCTGTGCA TCCACAACAT GTGGGGTTCT
92151 GGTCTGCCTT TCCACTATGC TGGATTAGTT TTGTATGCTG TGTAACAAAT
92201 TCCTACAGTC CCAGTGACCA GAAAGAACAT ACCTTTATCA GCTCGCAGTT
92251 TCTTTGGGAC AGGTGTCTGG GCACAGTCTA GTTGAGTTCT CGGCACAGCT
92301 GCCATTAAGA TGTCAGCCAG AACTGGGTTC TCTTCTGGAG GCTGAACTGG
92351 GCAAGAATCC ACTTCCAAGC TCAGTCAGAA TGTTGGCAGG AGGTATTTCC
92401 TTGTGGCTGT AGGACCCATG GTGGCTACTT TCTTTAAATT TAACAAGGAG
92451 AAGAATACCG TAGAGTAAGT TGGCTAGAAA GAAAACAGAG TACACATACT
92501 TGAATGATGA TATATAACAT TGTAACATAA CTCAGTCACA GAAGTAAGAC
92551 CATCACATCT GCCATGTAAT GTCGGTTAGA AACAAACCAT GGAACCAGCC
92601 CATGCTGAGG GGCTGGAAAT TATGCAAGGG TGTGAACACC AAAAGCTGGG
92651 AATCCTGGGG GTCACCGTAC ACAGTCTGTT CACATTTCCT CTAAAGAAGT
92701 TGCACTGCAT CACAGTTCCA TACCAATTTC TGCTATGACC TTAAATATAG
92751 CCCTGAACTT CCCTGTCAAG GAAGAAGTGA GGAGGTTTCA ACAAGTGATC
92801 AGTAATGATT CTTTTATGTC TAAGATTCTA GGATGATTTC CTCTCTGCCC
92851 TGGTAGGCTG CTCTTCAAAG TATGACCTCC TCATTGTTTC TCTGCTCTAC
92901 CACACACTCA TTCCCCTCCA AGAAGGCTGC CCACCTGTAA TGACCTGTCT
92951 ACAGAGCCTG TGATAGTGAC TTGTGATAAA TGGCTATTAG CACATTTACC
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FIGURE 3DD

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93001 AATCAAGGTC CTGTTTGCAA TTCGGTTGTG GGTCAAAATT ATGTTTGTTT
93051 TAACTGAGGT CTTTAGTTTA TTTCAGGCAG AGATCTGGGC TGGAGTGTCA
93101 CCTTTGTGTC TAATTCTCAC ACACTGTACT ATCTTAGCAG TCACATTTTA
93151 TTTTCTTGAG ATGATAATTT ATAGGAAAAA ATAAGACATT TCTGCAGCTA
93201 ATCATTTTAG TCAATGATCA TTGAGTGACA GGTGAGCTCC TAATAAATAA
93251 ATTTGCCAAC ACAGTGACAC CTCAGGTTTC TGAAGCCTGT GGGAATGAGT
93301 CATCTGGAAA GATGTTTTTC TAATTCCTGG AAGTATTTCA GAGATTTTTA
93351 ACTATTTAAT TTATACTACA AAGCACCTAT GTCACTTTTT TAATGACTTA
93401 ATAGGAGCTA TCACTTATTG TTTACACCAA GAACTGCGTA CTGTGCTAAT
93451 TGGCAGGTTC CACACACCAC CTAACTTGAT AATCAACAAT TCTCTGAGGG
93501 GATTAAGCAA CTTGCCAATA TACAGTCAGT ATATGGGGAC CAGATTCAAA
93551 TGTAGAATTA CCTTCTTCAA AGGCCCTGTT CTAGGTATAG ACGCTCTTAC
93601 TTTCACTCTT ATAATAATAA GATATCCTCA AGGTCAGATG AGCTGTTCAG
93651 TGCTGTTTAC CAAATAGCAT AAAACTTCAG TTTAGATACA TATTTTAGTG
93701 GGTAGGTACT ATATGTTAAT TTGTGCTCCC TCAGAAAGAT TTGTTGAAGT
93751 CCTAACCTCC AGTGCCTCAG ACTGTCATCT TTTTTGGAAA GAGGGTTTTT
93801 ACCCAGATAA TCAAGTTAGA ATGAGGCCAT TAGTGTAGGC CCTAATCCAG
93851 TATGACTGGT GTCCTTATGA AAAGAGGAAC TTTGGACACA GAGGAACATA
93901 CAAAGAGTGA AGATGATGTG GATGTAGAGA GACACAGGGA GGATGACAGG
93951 TGAAGATGGG GGATTGATGT GATGGGTCCA CCAGCCAAGG AATGCCAGAG
94001 ATTGCCAGCA AACCCACAGA AGCTGGAAGA GGCCTGGGAG GAGTCTCCCT
94051 GAGAAGTTTC AGAGGGAGCA TGGGCCCTGC TGGCATCTTG ATTTTGGACT
94101 TTCTACCTTC AGAACTGTGA GAAAATTAAT TTCTGTGTTC TTCAAGCCAC
94151 TGTTTGTGGT ACTTTGTGAC AGCAGCTCTA ACAAATGAAT GTAGTAAATA
94201 TGTTTCTATT GTTTTCTTTG CTGCTAATTT TTTAATCTTT GCTTCTCTAG
94251 TAGGTGCTAC TCAGAGCACC TTCTGTCCTC ACTCCTAACA TGCTGCTTAC
94301 AATACATTAT GGGATAGAAG ACCAAGTGAC AAAACTTGTT TGTATTGTTT
94351 GTAAAATTAA ACTAAACCAA GAGAATATTC AGTAAGTCAA GTCCATTGGC
94401 TTTAGTATAG GGTAACCTAT TTTAATGTTG CCAGAGACTG TCTTTGCTTA
94451 CTTTTGTATT TCAGGTTTGG GAAGATATTT TCAGTATCTG TAGGCTTTTT
94501 TTTTTTTTAT ACCACTTCTC CTGTCCAAGG TGTGTTGTTT TGCTTTTATA
94551 TATCTATTAG GAAAGTTAAA TCTTTTCCAT TTTACCAAAG CTACATGTCC
94601 AGTATGAGAA CATTTAAAGT CTAAAAATTA TCTGATTACT TATATTGTAT
94651 GTGTTCTGCT TGATGCTGGC TTTCTTTCAG TGTATTGATA AAAGTTTCTA
94701 TTTGTTGCAG TGGAATAATA GACTTTGGTT TTAGGCTATC ATCTGTGGAG
94751 TGCTTAAGAA AATGCCCTTT CTTTTTGTTT TGGTAAATCT TCTTTTCAGT
94801 AGACCACAAG CCCTTGCAAA TGTTCTCTTT TTCTAACTCT GGTAGCAGAA
94851 GGACCACTTG AGCCTCAAAA CAAAACGGCA GTGCAGTAAT GAGGGTATTA
94901 GGTTGATGTG TTCTATTCAG CACCTGCTCC CGAGCTACCG AATAATGAAT
94951 GAGCATGAAT TACACATTGT GAAAACAGGA GAATCTGCCT TCTTTGTGTT
95001 GTATGCATCA AGCAGTTTCA AAAGGGCTTT GCAATTGTGT TTCTCACACA
95051 AAGCCACCCA TTTGTGAAAA CCCATGTGTA AAGGCAAAGA GAACTGTCTG
95101 TGTACAGGTT AACATTTAAC TAGACTGGCA GAGCTTTTAA TAATTTCTAT
95151 AAGGTTAATG GCTTCGTTAA TATGCAACCT GTGATTTGGT CCAAGTTAAA
95201 TTTTACTTTG CCCAGAATAC ATTATAATAT AAAGCTTAAG CTTTATTCTT
95251 TCAGGTTTAG TCATTTAACA CATAATATTG ATCAATTATG CATGTTGGAC
95301 ACAGAGCTCT GAATAGAGCT TTGAAATATA AAACTATGGT TTTAGTCCTC
95351 TTAGAGCTAT GATGTTTGGT AGGTTAGGTG AAGTAGACAC ATTTTTGACT
95401 TATAAATTTT CAGCTTACAA TGGGTTTATC AGGGCGTAAC CCATTGCAAG
95451 TTGGGAGCAT CTGTACGATG GTATAGATAT ATATAATGCA TATAGTTTTA
95501 TATCCTTTTA AGACAAAATA TGAAGATATT TTATTTGCTC AAATCTTGTT
95551 ACACAGTTTT CCACTGTGAT ATTCACATGC TGACAGAGAG GCTATTTGCA
95601 TGGTGTTTGT CACCAGCAAT GAACAGCAGC ATTTGAGTTA TGTAGTGGCT
95651 CTGCCAGTTA CCAGTGGGGC AACTTGGGCA AGACACTAAG CACCTCTGAA
95701 CCTCATTTGT TTTATCAGTA AAATGAAGAT AGCTATACAT ACTTCACAGG
95751 CTGTGGTGAT GATATATTCT AATGAATATA CAGTCTTAAA TAAAAACATT
95801 CAATAAATTC TAGCTACTCA TTTATATTAA TTTATTATAC CCATTTGCTT
95851 TGAGTTATCT TCTTTGCAAT AAGCTGTGGG AAAAACTTAC TGTTCCTTCT
95901 CATACTCCAG GATACATCAT CACCCAAATC ATTACACATT CTTATATAAC
95951 GCAAACATTA AGAAAGAACA ATAATCTTAC TAAAAAGCAG AGTGTGGTAT
96001 GGTAGAGAGA TTAAGAGGCT TTGGAATAGT TACATCAGGG ATCAATTAGT
96051 GAGCTGTGTG ACTTTAGGCA AATTAATAAA CTGAATTTCT TTAAATTTTG
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FIGURE 3EE

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96101 TTAAATAGGT ATAATAACAT TATATATAAG AAAGCAGGAA AAATATGAAC
96151 AGCTCCTATT ATAATGCTTG CAAAATCAGG AGTGCTTAAT AAATGGAAGC
96201 CACACTGCGA TTTTCCAGAT AATTGTGAAA CAACTACGGG CCATTACAAA
96251 ACCATAGGAA ATTAGAAGTG AGGAGTAATT TGGAGACTGA CAAGCTCTAC
96301 CTTCATCTAA AGGCAGAATT TCTTCTGCAG TCTCCCTAAC AAGGAATCGT
96351 TATACCTCAG GGATGGGATA GTCACTACCA CATAAAGTAG TTCATTTTCA
96401 GACATGCATA ACCTTAGAAA GTTCTTCTCT TGATTTACAA TTAGCCTCAT
96451 AGTTCTGTTG CTGCCTATTG GAGTTTTACT ACGTGTACAG TCAGGCAGGG
96501 CTTCCATTCA GTCACCACCC ATTAGTACTG TTGTACTAGT AATTTATGGA
96551 TGGCGTCCAT TCTTACTGGT CCATGTCCCA TTCTGATTTG TGTTTGTGCC
96601 ATTTTTAAGT GTTTTGAATA TTAACCCTGG TATCAGATAA ACATGGAGTC
96651 CTGACTTTTT CCATAATCAT GAATAACAGT GGAATAGTTA CATCAGATTT
96701 GTGTGCCACT GTGGTCCCAT CTATGAAATA GGGATAATAA TTGTACCTAG
96751 TTCATAAGGT TGTTTGAGGA TAGTGTGGAA TAAAGTATAA AAAGGGCTTA
96801 GCCTGGTTTC TCAAATATTG CAATAAATGA AACTTAGCAT CATGATGCTG
96851 TCACAATGGT TCAATGATAA TTGAAAACAT CGATTCATCA TTTAGCATCC
96901 TCAGCTTATC AGTTTCTCAC TATCTAGCTC TTCTTACACT GGACACTTCC
96951 TAATTATTCT TTCAATGTTT TCTGGAAGTT AGTTGAATAA TTACTGTGCA
97001 CCAGATACTA CACAGTAGTC CCCCTTGATG CATGAGGGAT ACATTCAAGA
97051 CCCCCAGTGG ATACCTGAAT ACGCAGATAT TTCCAAACCC ATATATACTA
97101 TGTTTTTTCC CTTTTGTACA TACCTATGGT AAAGTTTGAT TCATAGAGTA
97151 AGAGATTAAC AATAACTAAT AATAGAACAA TTATAACAAT ATGCAGAGTA
97201 AAAGTATGTG AATGCAGTCC CTCTCTCAAA GCATCTGATT GTACCGTACT
97251 TACCTATTTT TGAACCACAG TTGACTGTGG GTAAAAAGGA AAACTGCAGA
97301 TAAGGGGGGA TTACTATACT ACGAGTTTTA CATGTACCAT TTAACTAAAT
97351 CATTACGACT CTATAAAGTA GATATGATTA TTGTCCTCAG TTACAAATGT
97401 GGAGGGCTGA GTCTCAGAAC GTTCTATTAC CGACATGGTT TTGGTCCCAA
97451 CAGAAAACCT CATAATGGTT TAAACAATAA AAGAGATTTA TTATCTTATA
97501 AAATCAGAAA ATCCAGATGT GTGCTGGACT TGGAGGGTAT CTTGATTCAA
97551 CAATTCAGCA GTATCACCAA CTAGCTGGTT TCTTTCACTC TCTTCTCTCT
97601 TTTCCATGTG GCCACTTCAT CCTCAGCTTG TTCCTCCATG TGATTGCAAG
97651 AAAGCTGCCT GCTGCCCAGG GCTCCATGCT AAATTCTTTA AATCTAAAGA
97701 ATCACACTCC TTCTCAAAAC TTTCCCCAGG ACAGCAAGGA AGCTTTTTCC
97751 TCAGAAGCCC AGAACATAAT TCTTTCTGAT ACTCAGTGGC TTAAATTGGG
97801 TCACCAGCCC ATCCCTGAAC CAATAACAGG GCCTGTGGGA TGGGATAACT
97851 CCTACTTAGG CCTGACTCAC ATAATCCTTC CCTACAGTCA GGGTGGAGTA
97901 GGTTTCCCAA AGCACACAAA ATACAGTGTG TGTGTGTATG TGTGTGTGTG
97951 TGCGCACGTG CATGCGTGCG CGTGTGTGCG CGCATGTGTG CATGAATGTG
98001 TGTGTTACAG AGAAGTGAAA ATACCCAGTT GAAAACTGAA ATGATGATTA
98051 AGAGAATGAA GAATGCGTAT TAGAAAGGCA ATCAAAATGA CCATTAGTAA
98101 GCTGCACAGT CGAGATCTGA GCCTTGGTCA TTTGACTACA GAATTAATAC
98151 TCTTAAACCT CCACTATCTA CTGCTTCCCA AATCAACCTA GAAATCCCTG
98201 GGGTTGGATA GGACCATTTG TGTTTGAGAC TATTACCAAC ATTACTAAGT
98251 ACTATACTAA TATACTCATG CAACCTAAAG CATATATATG TGAAGTGTGT
98301 ATATGTACCC ATATATATAC ATACACACTC ATATACTACA CACAGTATAG
98351 CCTATACAGG GCTCATGTTT AATCAGCATA CACTGGTCTG GCCCTATCAG
98401 TTGTATTTCA GTGTATTGGC TGATGAAGAG GTCATGCCTA AGCTTTGCTG
98451 CTACTCCAGC CCCTTTTCCA ATCTCCCCCT CATCCCCCAC CCCTTCCCTC
98501 CCTTGACCCA GCAACTGAAG TGCTAACTCC TGGCCCAGGA GAGGTCCTTC
98551 AGGGCACTGC TCCTGGGCTT CCATCAGCAT CCCTTCTGAT GAAAGGATGA
98601 CTGTGCTGTT CTGGTTGTTA AATATTTTGT CCATCACCTC TGGCTATTTG
98651 TAAATATATA TACTTACATG GAATACTATA TATGCCCACT ATATTTCAGT
98701 AAACTTTACT ATGCTAAGCT CTAGAGAGTT TAGATCATTT GTCCAAGATT
98751 ACATAATGAG TGACTGGGAT TACAACCAAA GATTGTGAAG TACAATCTTA
98801 GGAGGATGAT ACCTAGTCTT TAATCATCTA ACCCTGACAG CCTTTCACTT
98851 CTGCCCCCTA TTCCAAACTG TTTTTCCTTA TAATTTTCCC TCACTCGCTC
98901 TTAACATGGG TCTGTTTTTT GAGACCAATA GCCCATCTGT GACACCCTAA
98951 ATAATATGTT ACAGAATTAT ATGTATAATA TTTTTCCCCT CTCCAGAACT
99001 TGGCGATGGC CCAATCTGAG AGACTGTTAT GTGGCAAATA ATTAAATACA
99051 AACTATGGAC CATCAAAAGG CCATGGGACA CTGAAGGAGT TGATTTTGGT
99101 TTCGATATAC CGATTTCCTT GTTTGCTATT TTCATGTACA TGTACCGGTA
99151 TAGGATTGCA GGGTGAGCAA CTTGACTCCA GGGGAGGCGC AATGAAGGGA
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FIGURE 3FF

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 99201 TGTAATTAGC CTGTTAACCC TGCTAATGTC TTGTAAAGTC ATTCAAGTGA
 99251 GAAGAGTAGA TACATCAATT CTTCCTTGGA TCCTGCCACA AGGAGCATTG
 99301 TATTTCCACT CTGCTATTTA TAGTTCTCAC AGCTGGAATC AGCTGGTTCA
 99351 GCAGGACATG GCTCTTTTTT ATTTAATCAA ACCAAGATGC AATGAAGAAT
 99401 TTCCAAAGTA TGCATCCTAG AATTTCCCTT TATCACCCCC AAAATTCCAT
 99451 AGTCCCTCTG AAATCATAGG CTCGTAACAG GCATAAATCA CTTCTTATTT
 99501 ATTACTCTTA CTCTAATACA TACACATACA CTTACTGGAA AGTCAAGTTT
 99551 CTTAGTTGGC CAATGGTAAA TGTGGCGCAT CTGGCACACA GGGTTTGTTT
 99601 GGGTTGTTTT GGGGGTGGGG ATTGGTTGTT TTGCTTTGTT TTGTTTTCTC
 99651 TTCTCTTCTT AGGGGAAAAA GACATGCAGG GCTTAGTATT CCAACAATTT
 99701 GAGAAACCAG GGGGCTGGGA TTCATTCATT TTTATGACAA ATAGTTACTC
 99751 GAGCACCTAC TTTATTCTTG GGTACTTTTA TGAGTCCAGG GGCTGCTGCA
 99801 TTGAACAATA CAGAAAAGAA GTCCTTTCAC TTAGAACTTA CGTCCTAGTG
 99851 GGGGTTGGGG GTTGGGGGTT GAGAGAATGA AGCATTCTTA CAAAGAATGT
 99901 TAAAAGCGAA CTATGGGCAG GAATTGAGGA TATGAGTTTT GATGTATAAA
 99951 GAAAAAGTGA CAAGGTCAAT AATTGGTGGT CTTAGTGTGA TAGATATGCC
100001 AGTTTGGAAA TTGTATTGAA TAAATGCTAG TCAGGGCTA GGCTGTAGTT
100051 ATGAAAAGGA GATGATTAAG GAAGTGAGAA TAAGGAAACT ATTGGTGTGG
100101 GACGGATGAA AAGATTATTG GAGGCAAGTC AAGGAACTGA GAGGCCAGGG
100151 TGTTAGATGG AGCATTCATG TAGACACTGA AGTCACCAAG AATAATAAAT
100201 AACAAGTAAG AGGGAATTCA TCATTAGCTA TCTGCTTATG ATATGGATGT
100251 GTTTTTGCTG TGTCCCCATC CAAATCTCAT CTTGAATTGT AGTTCCCATA
100301 ATCTCCATTT GTCATAGGAA GAATGCAGTA GGAGTTAATT GAGTCATGGG
100351 GGTGGGTTTT TCCAATGCTG TTCTTGTGAT AGTGGGTGAG TCTCATGAGA
100401 TATGATGGTT TTATAAAGGG CAATTCCCCT GCACATGGTC TCTTGCCTGC
100451 CTCCACGTAA GAGGTGCCTT TGCTTCTCCA TCACCTTCTG CCATGATTGT
100501 GAGGGCTCCC CAGCCATGTG GAACTGTGAG TCTGTTAAAC CTCTTTTTCT
100551 TTATAAATTA CCCAGTCTTG GGTATGTCTT TATTAGCAGT GTGAGAATAG
100601 ACTAATAAAG CCAATTGGTA TGAGGAGTGG GGCACTGCTG TAAAGATACC
100651 CAAAAATGTG GAAGCAACTT TGGAACTGGG TAACAGGCAG GGGTTGGAAC
100701 AGTTTGGAGG GCTCAGAAGA AGATAGGAAA ATGTGGGAAA GTGTGGAACT
100751 TCCTAGAGAC TTGTTGAATG GCTTTGACCA AAATGCTGAT AGTGATATGA
100801 ATGAAAAAGT CCAGGCTGAG GTGGCCTCAT GTGGAGATAA GGAACTTACC
100851 AGGAACTAGA GCAAAAGTGA TTCCTGCTGT GCTTTAGCAA AGAGACTGGT
100901 GACATTTTTC CCCTGCCATA GAGATCTGTG TAACTTTGAA CTTGAGAGAG
100951 ATAATTTAGG GTATCTGATG GAAGAAATTT CTAAACAGCA AAGCATTCAA
101001 GAGGTGACGT GGGTGCTCTT AAAAACATTA AGTTTTATTC ATTCACAAAG
101051 ATATGGTTTG GAATTAGAAC TCATGTTTTA AAGAAAAGCA GGGAATAAAA
101101 GTTCAGAAAA TTTATAGCCT GATGATGGAA TAGAAAAGAA AAACCTATTT
101151 TCTGAGGAGA AATTCAAACT GGCTGCGGAA ATTTGCATCA GTAATGAGGA
101201 GCAAAATGTT AATGGCCAAG ACGATGGGGA AAATGTCTCC AGGGCATGTC
101251 AGAGGTAGCC CCTCCTATCA CAAGCCCTGA GTCCTGGGAG GAAAAATGGT
101301 TTCATGGGCT GGGCCCAGGG CCTTGCTGCT TTCGTAGTCT CAGGACTTGC
101351 TGCCCTGCAT CCCAGCTGTT TCTAAAGGGG CCAACATACA GTTCAGACCA
101401 TTGCTTCAGA GGGTGTAAGC AGCAAGCCTT GGTGGCTTAC GCATGGTGTT
101451 GGGCCTGTGG ATGCACAGAA GTCAAGAATT GAGGTTTGGG AACCTCTGCC
101501 TGGATTTCAG AGGATGTATG GAAATGCCTA GATGTCCCGA CAGAGTTGTG
101551 CTACATGGGC AGAGCCCTTA TGGAGAACCT CTGCTAGGGC AGCGTGGAAG
101601 GGAAATATGG GGTGGGAACC CACACACAGA GTTCCCACTA GGGCACCACC
101651 TAGTGGAGCT GTGAGAAGAA GGTCACCATC TTCCAGACAC CAGAATGGTA
101701 GCTCCACCAA CAGTTTGCAC CATGTGCCTG GAAAAGCTGC AGACATACAA
101751 TGCCAGCCAA TGAACGCAGC CAGGAAGGGG GCTGCACCCT GGAAAGCCAC
101801 AGAGGTGGAG CTGCCCAAGG TTGTGGGAGC CCACATGTTA CATCAGCGTG
101851 ACCTGGATGT GAGACATGGA GTCAAAGATT ATTTTGGAGC TTTAAGATTA
101901 TACTGCCCTG CTGGATTTCA GACTTGCATG AGGCCTGTAG CCACTTTGTT
101951 TTGGCCAATT CCTCTTATTT GGAATGAGTG TATTTACCCA CTGCCTGTAA
102001 CCCCATTGTA TCTAAGAAGT AACTAACTTA CTTTTGATTT TACAGGCTCA
102051 TAGGCAGAAG GGACTTGCCT TGTCTTAGAT GAGACATTGG ACTGTGGACT
102101 TTTGAGTTAT TGCTGAAATG AGTTAAGACT TTGGGGAATT CCCAGAACTG
102151 AGGGTTCCTC CCCATTGTAG ACCATATAGG TAGCTTCCAG ACGTTGCCAA
102201 GGCATTTGTA AACTGTCATG GTGCTAGTGA GAGTGTCTTT TAGCATGCTC
102251 ATGTATTATA ATTAGTGTAT AATGAGCAGT GAGGATGACC AGAGATCACT
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FIGURE 3GG

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102301 TTTGTCACCA TCTTGGTTTT GGCCAGCTTC TTCACTGCAT CTTATTTCTA
102351 TCAGTGGGGT CTTTGTGACC TGTACCTTGC AAAAACAGTC CTGCTGATTA
102401 CTAAATTCCT ATCTCACCTA TTCAAGATGG AGTCACTCTG GTCTGAATGC
102451 CCCTGATAAG AGAATCCACA GTGTTCAATT CTCCCCAGTT GATTCTGAAG
102501 CATATCCAGG TTTATTAGCC ACTAAGTAAA AATATATTAT AGACTACTGT
102551 CAATGAAAGA AACATTTTGT AAGTTATTTC ATATTTATTT TTACTTGAGA
102601 AGACTGAAAA GGTAAAGAAG TGATGCTAAA ATTTAGAACT AGAAAATCTC
102651 AACTTGCTCT AGTAGGAATT TTAATAGAGC ACACTAAGTT TCTTTTCATT
102701 TTCTCTCTCC TGGTATGTGA ATAAACAACC TTCCATACTG CAATTTACCC
102751 TGTAGTGAAT TAGATGTTAC CCTATTATAT TTTGGAGAAA CTATATAGTT
102801 AGAATCTAAG CTTAGATAAC TTATTTTTAT GTTACAAAT CCACTTTCTC
102851 TTATACATTT TTCTTAAATT TTTCTCATAT TCTTTCTCTG AATTTGTGGT
102901 AAAAATACCC CTTTCCCATT CTATGTCATG GTTCTTTACG AAGCTTTCTC
102951 ATCCTCTCCA TCCCGAGGGA ACTATGTCTC ATTTATCTTT AGGTTTTCTG
103001 TATCTTACTA CAGTGACTTA CCAGAGTAGG TAAATATCTG ATGAATAAAT
103051 GAATACAAGA TTTAATTAAG AAGTAATCAC ATTAAACTAA TTGTTCCCTC
103101 TCTGATCTCT GTAATATTAA GTTTCAAAGT AGTTTCTGGG AAAAGTAGTT
103151 AACACAATGA TGTATGGATT CAATAAATAA GAAAAATGGT GCTCAGGGAT
103201 TTAACAGAAA GCTCATAAAA TGTCAAATCC ACAGCAATTA ATTTCTCCCA
103251 GTAAGTCCTC ATAAATTCAG GCCAAGAAAT TTGATACTGA TCTTGCCTCT
103301 CTCAACTCTC ATCCATCTTT GGTAGGGCTC CTCTGGGCCT CTTTTTCACC
103351 TGGCAAACAG TACCTGATAC TCATTGGATG CAGATCTGAA AGAGGTGGAA
103401 AGAGCCCGAC ACCTGGTTTA TCTCTAGCTT TATGGTGCAG AGAGTATTTG
103451 ATGGTGTGCA CAGTGCTCTG TATATACTGT TAGGATCAGC CTTCTTGAGT
103501 GCACTGGAAT TTCTCTGGGT GTCATTAAGT TCTTCATTTA CTGACCATGA
103551 GGCACTGGGA TAGAATATGA TATTAATCAA GAAACCATCC CTGACATCAT
103601 GATCCACTTG GAAAACTTGC AGAAATTAGA AAAATTTTTT GAGTAGGCAT
103651 TTTGCTTTGT TGCCCAGGCT GGAGTGCAAT GGCTAGTCAG GGCACAGTTG
103701 TGCAATGCAG CCTCAAACTC CTGGGCTCAG GTGATATCCC TCTTCCACCT
103751 CCTGAGTGGC TGGGACTATA AGTACACACC ACTGTGCCTG GCAAGAATTT
103801 TTTTTTTTAG GATGTTATAA GGCCTATAGT TATTTAATTA TTAATCCTGG
103851 GGTAGTTAGT GAAAAGATTT GGACCAGTCT TTTACACACT GATGTACAGC
103901 AAGATAACTA TAGTTAGTAA CATTGTATTA TATACCAGAA ATTTGCTATA
103951 TCAAAGTATC ATGTTGGCCA CTTCAAACAC ACAATTTTTG GTTTAAAATG
104001 ACTAAAAAAA TTAAAATAGC AAAGTAAAAA AAATTCACAG GAGAGCACAA
104051 AACCCACCTT CTTCCAATGA AGGGAGTAGT CTGGTGGTTA ATACTTGGAG
104101 GATAGAATGA TAGAGTTTGC AAAGCCTTGG TGAATATTAT AGTAAGGAAC
104151 ACTCCTGAAT CAAAAAATCG CATTGTACTT TATAACAGCC CTCACTTTTC
104201 CACTCTCAGA TTTTTACTGC CTTTCCCTAA TGTACCATTA AAGCCCTTCA
104251 GCCTAAATTC ATAGACTCCA TTAGAGAAGA AATTCTGAAA CAGGTTTTGG
104301 GAACACATTC TCAGCCTAGT CAAATAGCTT TCATGCTGCT AGAATAAAAA
104351 TACCTTAATC TTTGACAGAC CAAGTCTGTC AGCTTACTCT TTACTTAAAA
104401 ATATTAATGA GTAACAAGTC CCATATCCAT AAACAGAACC AAGTGTGTGA
104451 TAAACTGTGA TAAATGTTAT GGTGGAAGAA GTATCCCATG TGGTCAGAAT
104501 ATATGGGATT AGGGGGGATT TGACCCAGAA ATGAAAAATC AGGAAGGCTT
104551 CCTGCAGGAA ATGGCATCTG AGCTGTGGGG TTAAGGGTGA ATCTGTGTTG
104601 TCTGAGTGCA CTGGTGAGAG GACTCTAATT TAGGCAAAGC AACAGCAGGT
104651 GTGGATGTGA GGAGGCAAAA GGAGACAGGG GGTGGTTATA TAACTACATT
104701 ATCAACCATA TTTTTCCCAT TTATAGTCTT TAAGCTCACA TCATCTGTGC
104751 AATTCTAGAG TTACACAAGA AAATGATGCT TAATACTACT AACATTACTT
104801 TATGGCAATG TAAATGCTTT ATATGATCCA ATGGACCAAT ATCTACATGC
104851 TTAGATACAA CATGCTATAG GAAGTTTAGA GTCTGAGTTT TTGAATGAGA
104901 GAGGCCTTGG TTCAGAGCCC ATTTCTTCCA TTTACTAGCC TGTGACCTTG
104951 GGTTAAGCTT CAGTTTTCTG ATTTAAAAAT TGGGGATTTT CTGTCTCATA
105001 AATTTACTGT GAGAATTGAA TGAGAAGATG AGTATTGAGA AGCTAGTACA
105051 CTGTTTCAAC TCCAGTTAGC TTTCTTAAGC CTTTTTGCCC CTACCCCTTA
105101 GTTCTGTTCG TTTTATTGTG AGCAACTTTC TTTTTTCTTT TTACTCCTCT
105151 AGGGATATGA AGCCTGACAA TATTTTACTT GACGAACATG GTAAGTGAGT
105201 GATTTGTTTG CAATCAAGTA CATGACATGC ATGTAGAAAA GTTGATTGTT
105251 CCCAGCAGAG GGGTATTACA CATGAAAAAG GTATTTTGTT CTATTCATTC
105301 GAGCTCTACT TACAAACTCC TCATAGACAA TATGGGGGAA CTTTATTACT
105351 TATGGCAGGT TATAGTACAA CAATACACCC TTAAATCACA TTGAATTTAC
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FIGURE 3HH

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105401  CTAATGAGAA  AATCATAGTC  TACTCAATTT  TCTTCCACTA  CTATATTTCT
105451  TCAAGAAAAC  CATCACAACT  TTTCAGTGTT  AGCTGGCCTT  AATATAACAC
105501  GCAATCACCT  ATTTTTTATA  ATGATACAGA  AGGCCTCAAG  CTGAGAGCAT
105551  TTGGCCAGCA  ATAGCATCTA  CCTAGACATT  AATGACATTA  TTTTGTTCTC
105601  ATTGCATCTA  CTTTTTTGCA  TTCCTTCTTA  TAAAAGGCAA  ATTGGTTTTA
105651  CATTTGCAAA  TTGGTTTTTA  CATTTACTTA  ATATCACAGA  AGAATTCTTA
105701  CATTTTAGGG  TCATTGTAAA  GACTGACCTA  ATACATGTAA  ACTACTTGAT
105751  GCAGTGACTG  TCACGAAGAA  ATCACTCAAT  AGAAGTCTAA  TATTGGTACA
105801  ATTTTTATGA  GGTGGTCATG  GGTTTCTCCC  CTTGGAAAGG  AAGCTGGAAC
105851  TGCTTCATCT  TGTTTTATGC  GGCTTTGTCT  ATGCTGGCAC  ATAACTAGTA
105901  TGTACCAATG  TATCTCAGAA  AAGATATCAA  GTTTTCTGTT  TAAAAATTTC
105951  AGTTTGAGAA  AAATCAGTTA  AAGAAAAACA  TAAAAAAGAT  AAAAGTATAT
106001  GTGTTATCTA  GATTTGTGAT  ATAGGGATAT  GGCAATAATC  AAGATGGTGA
106051  TAAGTGAATG  CTGAATTTCA  AGAACTACTG  ATTACACCCT  CTAGAATAAG
106101  CTTTTGCCCG  TGATGATTAA  ATGTGTACGA  TTTCTTCCTA  ATATTTATTT
106151  TTGTGTATAT  TGGGATTTAT  TAGAATATCA  GGGAAGATCT  GCAGGGCACA
106201  AAAACTGTAT  GTTATAAATG  TTAACAGTGT  CAATAAGATC  TTTGTTATGT
106251  CTTTAGAAGG  CTGCTAGATG  AGGAGAGTCC  TAGATCTTAA  AGGCTCCTTA
106301  TTCAATTTTT  ACAAAAAGGA  TTTGCAAGTG  GAACTGAAAC  TCCAAGTACC
106351  ATCTATTGCT  CATTATTTAT  TTACCTATTT  TTGAGCCTGA  TTTTCCTGAT
106401  CCCACCTGTG  CTCAGGGGGC  TAAGAAACAC  TGGTAATGAC  CTCTAATTTC
106451  AAAGCTCACT  GTCATTACTT  ATTTATGGAC  TGTCCAAAAA  GATTTTTTCC
106501  ACTTTCTTCC  AATGCCTTAT  TTCTTCCTTA  CCTTTACTGC  TTCTGACATT
106551  TGAAAACAGG  GTCTCTGATT  CTCAGAAATG  TGAGCAATGG  TGAGATTTAG
106601  CATGAAGGTG  ACTTTCTTTA  AAATACCAGC  TATCCAGAGC  TAGGTACAGT
106651  GGCAGGCACC  TGTAGTATCA  GCTACTTGGG  AGGCTGAGGC  AGGAGGATCG
106701  CTTGAGCCCA  GGAGTTTGAA  TCCAGCCTGG  GCAGCACAGA  GAGACCCTGT
106751  TTCTTGTTGG  GGGAAAAACA  ATTACCACTG  GCTTCTCTTC  TAGCCTATAG
106801  AGGCCACCTT  TGTGCAACTT  AGGGAGAAGT  GCTCCCCCTG  CCCACCACAG
106851  CTTCCTGACA  GCACATGGCC  CACCAAGGAG  AACCCAAGTT  AGGATTGAGT
106901  CCTCACTTGC  TCCCTCAGCT  GGGTGCCTTT  GTGCATGATT  TCTGCTGTTC
106951  CACCATTTAT  AGAGGCCTTA  AATGAAGGCA  TATAGGTCCT  ATCAATCCAA
107001  CACTTTCCCA  GCTTTATCCT  CCCTTCAGAG  AACAGTGTTT  TCATCCCAGG
107051  TCTCATCCAT  GGCTTCACCC  TACTTCTATC  ATTAAGGCAT  CCTATTCTCC
107101  TTCAGTCAAC  TTCTTCCTCC  TCCTCATTTT  CTTGGTGACT  TGGTCATTGC
107151  AGATGAGGAA  AAACATGAAG  AAATCAATTA  ATCTTCAAGT  TTAACCACCT
107201  TTAGAGACTA  CCCTTGTGAA  AGATTAATTG  TGTAACAGTG  TGGTTAAGAA
107251  TGTGACTTCT  GGAGCCAGAT  TGCCTTCATT  CAAAACACAC  TTCACTCATT
107301  TCCTAGCCCC  GAGAGCTTTG  ACAAGTTGCC  TAAACTTTGT  CTTAGTTTTT
107351  CCAGGGATCA  AAAGAATACT  TACTTAGAAA  AAAAATCTTA  CTTACAAAAG
107401  AAATCTTACA  GGGATCAAAA  GAATACTTAA  TTAGGGTCAT  TGTAAAGACT
107451  GACCTGATAC  GTGTGAAGTA  CTTGATGCAA  TGACTGTCAC  AAAGAAATCA
107501  CTCAATAAAA  GTCTAATATT  AGTACAATTC  TTCTGAGGCA  GTCATGGCTT
107551  TCTTTCCTTG  GAAAGGAAGC  TGGGACTGCT  TCATCTTGTT  TTATGTTTCT
107601  TTGTCTATGC  TAACACATAC  CTAATACGTA  CCAAATCTCT  ACCAGATAGA
107651  ATCTGTAAAA  GTTGTCCTTC  CCAAATAATT  ATTTTGATTT  AAGAAGTGAT
107701  ATACCAAATA  TTCTGCTTGT  CTACTTCTTA  GATCTTGTGT  TTAAACCATT
107751  TTGTTTATCC  CTTCATCCTC  AGGTAACTAC  ACTTTCCGTG  TACATTCTGC
107801  TGTCTTTCAT  GTGTGCAGGG  GGCAAGGGTG  CAGTCATGAC  ATTTTATTCT
107851  TGGTGGAGCT  GGGGCTCTGT  TGCCTACAGA  ATACAAGCCA  TCATTCCAGT
107901  GTGCCAGAGA  GAGAGTCTCA  GTCTGCCCCT  ATTACCTGGT  GTCTTATTTA
107951  CAATGACTGC  TTTCATTCTC  AAGGCTTTTT  AAAATTTGGT  CAGTGAATTA
108001  AGAAGAGGCT  TTTCTGTATT  ATATTCCTAC  CCTGAACTCA  ACTTGAAAAT
108051  CAATTGCTTT  GGGAAGGATT  GTATATGAAT  GGTACAGAAG  TGAGCAAACA
108101  AAAAAGACTG  AGAGCCATTT  TCTAAACATT  GCCTTAGGGA  TCTCTTTCTG
108151  GAGATAATAA  TTTTTTTGAA  GTTATTTACT  TCGTTTGTTC  AGATTCTGAA
108201  AAAGTAGGAC  TCTCAGACAT  TACTCAAGGA  ACATAATTAA  CCACTTTTCC
108251  ATGAACAAAT  TCCTGTTGTT  CACCTCTCCC  CAGCTCGTTA  TGTAGAGCTG
108301  ATCTTGTGAG  AATCAGCTGA  ATCACAAATC  AATGCCTGCC  TTTTAGAGTG
108351  TCTGCTGGTG  TGACTTTCCA  TGTGGAGCTC  ATATTTGAAG  ACCTCATTTG
108401  CCTTCTCCAT  CTCCATTTAT  AATATTTCAT  CCCTGATGGG  CTGTCGCTTG
108451  GGCCTCATGT  GGAAATTGTA  GCCACTGTGA  AGGGTAACCA  CCTATCTCTC
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FIGURE 3II

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108501 TGGTGCCCCC TATGCGCATC CCTACAAGTG AGCTGTGTAT CACACCATGC
108551 TGCTTACATT TTTATGCAAC ACGATTCAGT AACAGGCAGA AACTTTTATT
108601 CTTACTGACT CATATTCTTT ATATTCATCT GAAAAGATTG ACATTTAAAG
108651 GAGCCAATTG TACAATGGGA AATCCACTGT GTGAATATTT CTTGTACATC
108701 AGAATTTGCC TTAAAAATGT TTTTAACTTA GAGCACATCT GTACTGTTCT
108751 CCCCAAATGT CCCATTTACT AGTTCAGAGC AAGATGACAT TAGGTCTTGG
108801 GTGACTCCTG ACCCACTATC CTAATGTATA TTTTCATTTC CTACCAATGT
108851 AAGTACCCCA TCCAATTCTA TCAATACCAT AGTGTCTAAA ATTCTTGTAT
108901 TTTTCTTATT CAGGAAATGC TACAACCAGA GGAACAGTAA TGTCTGCCTG
108951 ACATATCAGA GAAAATGACA ATTATGTCAT CATCTGTCAC TTAGGTTTCT
109001 TAATACCATC CTGTTACAAG GAATAGAGGC AAAAACTCAG CGTAGGAGGT
109051 GAGAAAAAAC TGAGGCTGCC ATCTTAACAG CCTTTTCATT GCAGAGTCTC
109101 AAAATGTACC AAAAGATGAA GTGGACAGTG TCCTTTTAAA ACAACATACA
109151 GTGTAGAATA CAGTAACTTA TCCCCATTTA ATTACTCCCT AGGTAGTGCC
109201 TAAGGATATA CATTTTCAGC AAGGATCTCA GAAAAATGTG GGGCACATAT
109251 TCTAAACACC TGCGAGTAGC AGAGACTTAA AAGTTGGGAG CAGTGCCAAC
109301 TGATTGGTTA TGGTGCCCTA GAGCACTGCG TTGATGAAAG AGATCCTCAG
109351 GCTGTGCACA GGAGCAGCAA GAAAGAGTGT AAATGATGAC AACAATGATG
109401 GCTGAATTCA ATGGCATCAT AAAATGAATT CAGATTTTTT ATATGATCCT
109451 CTATCCCAAG CAATAGAGGC AAAAAAAAAA AGGCAGAAAC CCTCTCCTAG
109501 AGTGGTAAAT TAGGAAGTTC TGAGGCTTGC ACCTGAAAAA CTTTTCACTA
109551 AAGTAGTGAT TCTCAACTGG GCGTAATTTT GCTCTACTCC TTCTCCCTGC
109601 AGAGGACATT TGGTAATTTC TGGAGACATT TTTGATTATC AGGATTCCAG
109651 CCAGGGTTGG GAGGTGATAT CAGCAGCTAG TGGGTAGAGG CCGGGATGCT
109701 AGCATGCATC CTGCAATGCA CAGGACAGTT CGCACTACAA AAAATTATCA
109751 GGTCCAATAT TTCAATGGTG CTGAGGTTGA GAAACTCTGC TCTAAGGCTC
109801 ACTCAAGGCC TGGGCTAATG AAAAAAGCCA GAGAAGTCCT TCATTCCCAA
109851 GGCAATTCCT GTGTCCTTCA GTCAGCAGGA GACTGAACCC TTTCCTGTGA
109901 TCCAGCAGTC AAATTTCATT TTCAAAACAC AGAAGGGAAC CTGGCAGATA
109951 GGTCACCATG GTAAGGAGAA GCAAGTCATG GCTGTAGCCG GACCTGGGAC
110001 TAAGGCTTAG GGCCAGCACT CTGTGAAGTT CTGCCTTCAT TGTTTAGCTC
110051 AGAAGCACCA GGTTACAAGA TCCAGTAGAA CCTGACCCTC AAATAATTTC
110101 TCCCTCTCCT TAAATAGGCA TCCTGGAAGT GGACTAGAAC TCTGAGCCAA
110151 TCAGAAATTA ACTGTTTTAG GTTATTCAGT TCTTTGATCT TGTGATACAG
110201 CACACAAAGT TTTTGGTAGA TTCATAGTCT GACAAAGGGA TTCTAGACAA
110251 AATTCTAGGT CTTAACTCCA GCTCTGTAAC TTTTGAGTCT TTTGAACCTA
110301 GCCATAAATG ACTCATATAT AAAATAGGGC CTACCTCACT AGGCTAAAGG
110351 AGAAATTTTG TGCAACAACA TTTTGAAAAC TGAATCATGC AAGTGTAAAC
110401 AGCATTTAAA AGGAAAATAC TCAACATTCT TTCAACTGAC GTGTAATGAG
110451 TACTCACCAG AGTTGAGATG TTCTGCTAAG CCAGGCCCTC TTTTAAAAAT
110501 GTAATCTCAA ACTTTATTAG GTCTCATAAT CACCTGGAAG GCTTATTTAA
110551 ATATTGGCGC CCAACCCACA GAGTTTCTGA TTTGTTATAA TAGAGTTGAG
110601 GGGGGACGGG GCGTAAGAAT CTGCATATCT AACAAGTTCC CAGGTGATGC
110651 TGATGCTGCT GATCTGGGCA CTACATTGTA GGAATCAATT GGCTCTAAAA
110701 CCTTCTCTAC CTTCCACTTC TACATGAGCA TACATAATCT TGTAGCTGAG
110751 TCAGCTTGGA AATCTATGCA GACTAAAGTA GACAGTTGCA TGTCTGGCTG
110801 CTCATCTGAA TCACCTGTGG AATTTGTTGT TTTTAATACA GATACCTGGC
110851 TCTCCTACAA GTCCCACTGA ATTGGAGTTT CAGGAGACCG AAGCCCAGGC
110901 ACATGTATTT TGCAAAACTA CACTGAAGTT TCTGATAATG ACGGATATCA
110951 ACAATTAAAC GCTTACTTCT TGCCAAATGC TGTGCTAAGT CTCCTGTAAT
111001 CATTCTTTCA TTTAATATTT CTAATAACCT CTTGAGAAGA CTATGATTAT
111051 CTTTCCAACT TTACAGAGAG GATAAGTGAC GTTTTCAAGG TAACACAGCT
111101 AGTTAGTGGT AGAACCTAGA CTTGAAGCCA AGCAGTCTGA CTCCAAGAAA
111151 CAGGCTCTTC ACCACAGTCT CCAGACTCAC CTGATTTGTA TTAAACTTTG
111201 TGAATCACTG ATCCAACACT ATGAGCAGGA CCCATGGGGA GAAAGAGAAA
111251 AAGAAAAAAC AGAGACAACC TACGCTATGA TAAAGTTATT GAAATCAGGC
111301 ATTGGTGCCA CTCCAGCAAG AATGAGTGGC TACCTTTTTT TTAGATGAGT
111351 GCTACCTTTA CTTTACTGAA ATATCATGAC ATAAACAAAG CCAAAACACT
111401 TTCTGCACAA AATAAAATCC TGGTGATAAA GGCAGTGGGA TTTATGCTTA
111451 GCAGCAGGCT GGATACTATC AGGGAGCAGA CAAAGAAGTT TGATACAGGG
111501 CTTGTGGACT GTGGGCCCTG GAAGAATCTG ATGACATGCC CTCCAATTAC
111551 AGCTGTATCT CATCAAAACC ACAGACACAT GTAAATGGAA ATGCCAACAC
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FIGURE 3JJ

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111601 TTCAAGATTC TCTGAAAGCA GTTGACTGTC ATGCCAACAG CTAACATAAT
111651 AGGCTTGTTT GCCTGAGCTT TTGGCACGGC CCTTTTGTTC CCTTTAGCTG
111701 TAAATGCAGG GACCCTAGAG CACCTCATAG AGTGTGTTCC CTGCCACGTA
111751 TAAGTATTAG ACCCACACTA TATTGCTTTG AGTGTTAAAG CTGAAAGAGA
111801 CCCTAGAGAT CATTTAGTCT ACTCCTTCTT TTTTTATGTG AAGGAAAATT
111851 TAGATCCACC TTGGAAAAGG ACTTAGAGTC TACTATGTGT TAGAGGCTGA
111901 GTTCAAGGCA GAACCCAGGC CTCCTGGCTC CCAGTCTAGT GCTCTTTATA
111951 GAATCCCTTT AAAAATGAAG TTGACTGGCC GGGCGCAGTG GCTCACGCCT
112001 GTAATCCCAA CACTTTCAGA GGCCGAGGCA AGCAGATCAC GAGGTCAAGA
112051 GATCGTAGAA CACCCTGACC AACATGGTGA AATCCCATCT CTACTAAAAA
112101 TACAAAAATT AGCTGAGCAT GGTTGTGCAT GCCTGTAATC CCAGCAACTC
112151 GGGAGGCTGA GGCAGGAGAA TCACTTGAAC CCGGGAGGCG GAGATTGCAG
112201 TGAGCCGAGA GCACACCATA ACACTCCAGC CTGGCAACAG AGTGAGACTC
112251 CACTTCAAAA AAAAAAAATT AAATTAAATT TAAAAAAAAC CTAAAGTTAA
112301 ACCCCGCCCC CCACCCACCG CCCCCCGCTA TCCCTTGATA ACAGTTATTT
112351 TGCTGGGAAC TGATGAGGCC AACCTGAATT ATCAGACAAA AAATATGTAC
112401 AAAAATATTT TAGAAAAACT GAAGAAAAGG GATGCTTTCT TGGCTAGGAA
112451 ATAAATATTT GTATCCATAT TCATGCCAGT TTTGTAGTAA TAATATTTGC
112501 CTCTTACTTT TCTTTTCTTT TTTTTTTGAG ATAGTCTCAC TCTGTCACCC
112551 AGGCTGGAGT GCAGTGGTGT GATCTCAGCT CACTGCAACC TCTGCCTCCC
112601 AGGTTCATGT GATTCTCCTG CCTCAGCCTC CCAAGTAGCT GGGATTACAG
112651 GCACCCATCA CCACGCCCAG CTAATTTTTT ATTTTTTATT TTTAGTAGAG
112701 ACAGGGTTTC ACCATTTTGG CCAGGCTGGT CTCGAACTCC TGACCTCAAG
112751 TGATCTGCCC ACCTCAGCCT TCCAAAGTGC TAGGATTACA GGGGTGAGCC
112801 ACCACGCCCA GCCTATTTGC CTCTTTAAAA AAAATAATCC CATAAGGGAT
112851 GTTTGGAAAC GTGATACTTT GAGTATCTCT TGGCTGTCTC CTTCATAGTA
112901 TTCATAGGCT AAAGTAACTT AAAATGTCAC CAACAGACAA AAGATGCCTA
112951 ACTAGAATTA CCTGACCACA AATTCTTAAC TACTAAGGGT AAAACTTTTC
113001 TGAGGCTGAA CTACAGGCTT ACAATCAGAG ACTAATCATT GCATATCATG
113051 AAATGGAGAA TTGTTGGTTT AAGACCATAT CGGCCTTGAG GATGGACTGC
113101 AACTGGCCTA CAAGAATTAA CAGACTAATT GGGTGTTTTC AGTTAAAAGC
113151 ATGATTGTGC CACTGGGTTG AATGGGACTT AACTTTCTGT GTGGTTCTTC
113201 TCTCTCTGCA GGGCACGTGC ACATCACAGA TTTCAACATT GCTGCGATGC
113251 TGCCCAGGGA GACACAGATT ACCACCATGG CTGGCACCAA GCCTTACATG
113301 GGTATGGGTT TCATGAGTGT CTTTTTTTTT TCTTTCCTGT AAATACCATT
113351 TATTACAGGT GGAATCATCT GTGGGGATTT GCAGCTAGAA CTGGTAAGTT
113401 CCTCTCTGAC TTTACCTGTG GAGCTTCTGA TTTCATGGGT CTTCTCCACT
113451 AGCAAGCACC CAAGATGACT TTGATAGGAA AGGACCATTG ATTACATTTT
113501 GAAAACTTAC TTCGTGTGTC AAGGAAGACC GTTTGTACCC ACTTCCTAAC
113551 AAAAATATTA ACTAATTCAA TAAATACCTA CTAACTGTCT CTGTGTGCTT
113601 AGCACTGTTT CAGATGCCGG TGACCCTGTA GAAAGCAACA CAGACAAGGT
113651 CTTCAGATCC TGGAGCTTAC ATTCTAGTGG GAGCAGATTT ATAAAAAAAA
113701 AAGAACCAAA CAAGGCCGGG CATGGTGGCT CACGCCTGTA ATCCCAGCAC
113751 TTTGGGAGGC TGAAGTAGGC AGATCATGAG GTCAAAAGAT TGAGACCATC
113801 CTGGCCAACA TGGTGAAACC CTGTCTCTAC TAAAAATACA AAAATTAGCT
113851 GGGTGTGGTA GCATGCGCCT GTAGTCCCAG CTACTCGGGG GGCTGAGGCA
113901 GGAGAATCGC TTGAATCTGG GAGGCGGAGG TTGCAGTGAG TCGAGATCGC
113951 GCCATTGCAC TCCAGCCTGG CGACAAAGCG AGATTTCGTC TCAAAACAAA
114001 CAAACAAACA AACAAACAAA GAAGTAGGAA ACAGTAATAA GCAAAATGAT
114051 AATAAGTGGC AAAGTATTAT TTTAACCATT ATTTACATAA TACTGCATTA
114101 CATACATAGA GCTATAAACT TTACAAAATA CATTCCCAGC TATAATTTTA
114151 GATTTACTTG TAGTGCCACA ACAATCCCAT GAATTCTTCT GTTTAAAGAT
114201 AAGGAAATTC TGGAGCTGGA TGGTGGCATG CATCTGTGGT CCCAGCTGCT
114251 TTGGAAGCCA AGGCAGGAGC ATTGCTCGAG TCCAGGAGTT GGAGGCTGCA
114301 GTGAGCTATG ATCATGCCAC TGTACTCCAG CCTGAGTGAT AAAGTGAGAC
114351 TCTGTCTCTA AAAACAAATA AATTATTTTT AAAAATAAAT AAAGGTGAGG
114401 AAATTCTGCC TCAGAAAGTT TAAATGTCTT TGCATTATTT TGTGTGTAGC
114451 GAGGTGAGGA ACTGGTTTTT GCCTTGACAA TTCAGCATTT ACTAAGGGGT
114501 GACCAAAAAG AGAGTGTTAG ATGCAAAATT GTCAGTTGGT TTCACGTATA
114551 GTTGTGGTAA CAAATCAACT ACAAAAACTC TAAGTTCACC TGTTGGGAGC
114601 AGCCATCTAT ATAGACACCA GAACTAGTTG TTAGCAGAAC CAGCTTTACT
114651 TCCCGTCCAG CCTCAACAAT GCAAGGAGAG AGCTAGTGTC CTCGAGGGGG
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FIGURE 3KK

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114701 CACACAGTAT TCAGAAAGAG GGAGTTCTCC CTCCCTTTTC CCTGTGGTTG
114751 CTCCTAAGGC AAGTGAGTCA GATCTCAAGA GAATTATCTG TAAACTCTTA
114801 GAGTGACTGC AAGAAAAGAT ACCTGGAATT TAATTCTTGA TTAGATATCT
114851 GTGTAGTTAC TGGACTTGTG ACTGGTCCTG GAGTTAACAC AGCCTGGTTG
114901 GCCATGGAAG TTTGATGAGT TTGGGGGCTA GTCTTTCTGG GGATCATAGC
114951 AGCAGGAGAC AGGTATGCAG TGAATGTGAT TTGTCTTGGG GAGAAGGGAG
115001 GTGGATTAGC TACAGGCTGT GATCCACCTT CACATGGGAC CCTCCAATGA
115051 CCAAGAATAT AGCCTGGAAG GGAGGGAGGC TCCTGTCAGT GTGACTTCCT
115101 GAAAACACCA CAAGTCCCAA TAGAGCTCAA CATATCAGAA TCACTGAGAG
115151 TGGAGTCTAG GCATAGTGTG ATTTAAAGCT CTTAGCGTAA TTCCTCCGTG
115201 TAGCTAGGAG TCACAACTTC CACCACAGAC CCCTAAAGAG AGATTACTCT
115251 GCAGGGTAGC ACATGTGTGA GGACCCCTCT GCCTCGACTA CCCTTCTTTC
115301 ATGTCCTAAA ACAAATAGTG CTTTCTAGGA AAAGATAGAA GGACGTGTGT
115351 GAGAGCCAGA TCAATCCTCC ACCTCCATAC CGGGGTGGCT GAAACCAGCC
115401 CAGCAGGGTG AGTGAAGGAG CTTTGAATCA GATATAAGAA TAGTTTTAAA
115451 ATTCACAGAA CTGAATTGTA AAGCATCTAA AGTAAATGTA ATAAGCAAAT
115501 AGGACTAAAA CTTATTAGGC AACAGACTGA GATATCATTA GGCGAGCTCC
115551 TTATCCAGCA AAAACAGGAA GTTAGACACT GCACAGTTGC TGTCAAATGA
115601 CAGAAGACTA AAAACTACTC ATGCTTGGCG GGGTGCGGTG GCTCACACCT
115651 GTAATCCCAG CACTTTGGGA GACCGAGGCA GGCGGATCAC AAGATCAAGA
115701 GATCGAGACC AGCCTGGCCA ACATGGTGAA ACCCCATCTC TACTAAAAAT
115751 ACAAAAATTA GCTGGGCATG GTGGCGTGCA TCTGTAGTCA TAGCTACTCG
115801 GGAGGCTGAG GCAGGAAAAT CACTTGAACC TGGGAGGCGG AGGTTGCAGT
115851 GAGCCGAGAC TGTGTCACTG CACTCCAGCC TGGCGACAGA GTGAGACTCC
115901 ATCTCAAAAG AAAAAACAAA CAACAACAAC AACAAAAAAC CTACTCATGC
115951 TTTACCCTAA TTAGTTAAGA TGCTTAAAGC AGGTGATGTG GTGATGTTGC
116001 TGTTTAAACT GGTGGGATTA AGTCGGGTGG AATGAATTGT TTCAGCTAGA
116051 TATGGTCAGA GTAATTCAAA GGTAAAATAT TTCAACTTGA AATCAAGGAC
116101 AAGAGCAATG CCATTTTCTT TTAATATTTC ATTCTCTTCC CCCATGTAAC
116151 TAGAGAGAGA GAGAGAGAGA GGAAAAGAGA ACCCCCTACA TGCAGAGCCA
116201 CCTCACTTTC CAACAGAAAT CTTCTATGAG AAAAAAAAAT GAGCCTTATT
116251 TTCTATGATA TTTGAACAAC TGCAAATTTC ATGGCTTTCA ATTACCAGTG
116301 GGGGGAATAA ATCTCTTTTG TCACTTCTAA AATAATGGAC ATATATAATT
116351 CAGCCTATTT TCTGCCTAAA ACCTATGGTA CTCAAATGAT AAAAAAGCAT
116401 ATCCAAGCCT GCTGCTCTGA TGAGTTTATT CTCCAGGTTT CCTGGGTTTC
116451 CATATTAAGG GCTATTTTCT TGGAACCAAA TCAGAAAATG TGCATCTGGG
116501 TTTCCAGGGT TGGTTTCCAT GGTGAGAGAA GTACGGGGAG GCCACCTTTC
116551 TTTCCTCTCC CCAGTGGTTT TAAGTACAAT ATCTGTATAA TGTAATTTTT
116601 TCAAAGTTGG CATTTCTAGT CTTCTCACAA GATAGAACTG GGAAATTGGA
116651 ACCTAGGAAA AATTCTGTGC ACCTTCCACT TTTACCCTTG TAATTAACAA
116701 TGACTAATAT TTCTTGAAAT CTTTCCCTGG ACCAGACAAG GTGTTAAATG
116751 TTTTACATTC ATTTATTTGT TTATTTTTCT CAGCAGCCCC ATGGGGTGGA
116801 CTATACTTAT CACTACTTTA TAATGAGAAA AATCAGAAGC TAAATAATTT
116851 GGCCGAGATC ACATGGCTAA TAATTGAAAA GTCTAGATTT AAATCAAGCT
116901 CTGTCTGATT TCAGAAATCA AGCTTTTTCT TAAAAGGAAG ATTAATGAGA
116951 AATAAAAATA TATATTTGTA AATATTTTTA TCTGTGGTTT TTAAATGGTT
117001 CTAAGTCAAC TTAGTTAGGC TAACATATTC GAAATGTTTC TTGCCTTATT
117051 CCAAAATGAT TATGTGATTG CCACACTCCT CCTTTTGGAT AGGAGTCTTT
117101 CCCAGACGTA TTGTGGGTAG AAGTCTGCTG TCTCTTTTTA AAAATTATGC
117151 TCCCAATGGT TTGGTAAAAT CTACCAAATC TATCAGCACC CATTTTATAG
117201 TGCTTTCATA GGATACTAAG TAGCAATTCA CCAGAAAGAA CAAAAAGAAT
117251 TCTAAAAAGA AAGAAAACTA ACCAAAATAC TGAATGAAGA TTGGAGAAAT
117301 ATTCATCTAC TAATACAAGA TGCTGAGCAT ATTTTAAATC AGTTCCATAG
117351 CTCTGTAAAT AATAAGACAG TATGCCAGTT CTTCACCACC TTCCATCAAG
117401 CAAGGAAAGT TTTGCTTTTT ACAATTTATT GTCCTCTACC TCTGTGCTCC
117451 CTCTGGTCCC TCCATTATTC CTTCTCTCTT CTCCTTTGTC TGTATGAATA
117501 TAATCCAGAT TACTTAGAGT TAACCAATTA AAACCTTCTC CGCCGGGCGC
117551 GGTGGCTCAC CCTGTAATCC CAGCACTTTG GGAGGCCGAG GCGGGCAGAT
117601 CACAAGGTCA GGAAATCGAG ATCATCCTGG CTAACACGGT GAAACCCCGT
117651 CTCTACTAAA AAAAATACAC AAAAAAATTA GCCGGGCGTG GTGGCAGGTG
117701 CCTGTAGTTC CAGCTACTCG GGAGGCTGAG GCAGGAGAAT GGCGCGAACC
117751 CGGGAGGCGG AGCTCGCAGT GAGCAGAGAT CGCGCCACTG CACTCCAGGC
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FIGURE 3LL

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117801 TGGGCGACAG AGCGAGATTC CGTCTCAAAA AAAATAAAAT GAATAAAATA
117851 AAAAATAAAA ATAAAAATAA AACATTCTCC TCCAAATTAT ATATGTATGT
117901 ATGTGTATAT ATGTATATGT ATGTGTGTGA GTGTGTGTGT GTATATATAT
117951 ATATATATAT AAATAAGTTC ACTATGGACT AGCAAGCAAA AGGAAAGTAA
118001 TAATCCCTTT GCCAATAGAT ATTTATGGTT TATTTCCAGA CATTTTTTCC
118051 TAAGCACAAA CACATACTGT TTACATTTTT TAAATATTCG ATCATGCTAA
118101 ATGTAACCTA AATTTTCATT TTATAATGTA ACAATAATGA TAGCATCATA
118151 TAGTGAACAT TTATTGTTCC AAGCACTTTG CTAAGTTTTT AACATTTATT
118201 ATTAAACTCT CAACCCCATA AAATAGGTTT TACTATTGTT TAGATTTTAC
118251 AAGTTAAAAA AAAATCAGGC CCAGAGAGAG AGAAAGTGAT GTGTTCATAA
118301 TCACACAGCC AGTGATTGGC AGAGCATGAA ATTAAACCCA AGTCTAGAAA
118351 CATGCCGTGC CTGAGACATG GACGATGATG TGACAATGAT GAAGGTAGAA
118401 TGTCTGACAT TGCTAAGCTC TTCCTAAATG TTAAGCACTG TTGTAACTGC
118451 ATGCATTGTC ATTTAAACTA AAAACAGTTC TGTGAGGCCA CTACTATCGT
118501 TACAGTTTTA TTATTGCATA ATATATTAAC ATATAATTAA TGTAGTATAT
118551 TGTATATATA GTACTATTGT TATAGTATAT ATTGTTCTCA CTTCAGAAAT
118601 TAGCAGACTG AAAGGTTAAG AAACTTGTTG ACTGTGAAGC TGGAGACAGT
118651 CATAGGGGTC TGATGCCAGA GCCCTAACTC TTAACATGCT GCAGTACTGT
118701 CCCTTTGTTC ATGTCAATAA ACATGCCTCT GCTAAAATAG AAACCCACTT
118751 CTCTTAATCA ATTTTTTATT GTTGAATGTT AGGTTGTTTC TCATTTTGAA
118801 ATACAGATAG AGCATCCCAA ATCCAAAATG CTCCAAAATC CAAAACATTT
118851 TGAACACCAA CATGACACTC AAAGGAAATG CTCATTGAAG TATTTTGGAT
118901 TGATTTGGGG ATTTGGGATG GCCAACCAGT ATAGTGCAAA TATTTCAAAA
118951 TCTGAAAAAA AAAATTGAAA TGCAGAACAC TTCTGGTCCC AAGTATTTCA
119001 AATAGGGGAT ACTCAACCTG TACATTTAAA TTTGTAGTAA AAATCCTGTT
119051 AGCAGAATTA TGTCCTGGAA CTTAGTTATT TCTTTGTGAT AAATTTTCAT
119101 TCAATAATAA TAGTGTATTC TCTTACTGAA AATCACTCAA AGAAAATTTT
119151 GTGTTCTCAC CACAGAAAAC AGTAATGTGG GTAATGTGAG GTAAGGCACA
119201 TGTTAATTAG CTCTATTCAG CCATTCTAAA ATGTATTTAT TTCAAAAAAT
119251 AGTGTCATAT ACAATATATG CAATTTTTAC TTCTTAATTA AAATTAATTA
119301 ATTTGATTAA TTAAAAGAGC AAAAGAATTT CTGGTCAAAG CCTTTACATG
119351 TTAATAGATT TCTGTTCTGA AAATTCATAT TAACTTGTAC TTGCTCTGGA
119401 AGTGTCTGAA GATATTCATT TCCCTGCATT CTTATCAGTG CTACACTATC
119451 AATATCTTTA ATTGTCCCAA AAAAGGTAGG TAAAAATGAT ATGACATTAT
119501 GATATTACCA CAGTATTTCT TTGACTTCTT TTGTCAATTG CCTGTTCAAA
119551 TTCTTTGCTC ATTTTCTATT AAGGTGTTAA TACTTTTATC CTATTCCAAT
119601 AGTTCTTATT GATTATATAA ATAATTCTTG CCTTTTATAT ATTTGGAATA
119651 TGAAATCCTA GGGTATCATA TTTGTTGTAC ATTTCATTAC AAATATAATT
119701 TCTCATTTTT AATTTGTTGC TGTTTTATGG CCTAGTTTTG ACATGAAAAG
119751 CTTGCTAAAA ATATTATCAA GCCACTCATC TTTTTACTTT GCTTTCTAAC
119801 TTTGATGCTT TTCTTAGCAA GACCTTCTTA CCAGATTTTA GATGTGTTTG
119851 CTTAATATTT TTATTCTGAT TATGGTTTCA TTTTTTTACT TAACTCAGTT
119901 GTATATTATT TTGACTGAAC GGATGTGGCA AGGATCTGAC TTTATTTTTG
119951 TATGATTATT AAATAATTGT TTTGAGACTA TGTATTAAAT AAGTCCCTTT
120001 CCATGCTGAT TTGAAATATG TTCATCATAA ACTAAATACA TTTTTGTGCT
120051 AATATCTATA TTCTGTAGAT TTCAAATCTT GTAGCTTTAT AGGTTAATAC
120101 ATGGGATGCG GGACTCTTTC TTTATTCTTT TCCAAAAATA TTACTTCCAC
120151 AATTTTTTTC TTGTAGATGA AATTTAGAAT CATTTTTGTA AAGTTCCATG
120201 AATTAATCCC ATTAAATGTA TAGATTAGTG TTGGGTCCCT TTCTTTATGT
120251 CCTGACCAAA ATTTAATACC CACGTTTAAA AAAATCTGAA AACCAAATGA
120301 TGGAAATCCA AATATTTAAT AAATATATTA AAATGTAGTC AAGCTTATTA
120351 GTAAACAAGA CAATGCCAAT TTAAACCACA GTGAAATACT ATTACACACT
120401 CACCAGATTG GCAATAAAGG GTCAGTTATT GCCAAGTGTG GGTAAGGATG
120451 TTCAACAAAA GGAACCCTGA TCTAATACTG GTCATAGTGT GAATTTATAC
120501 AACACTTTGG TAAATAGTTT GGAGTTACTG TGGTACACAG AAAAGTTACA
120551 CATTCTTATC ACCAACAGTT CCCCTGCCAG GAATACACTC TAAAGAGATA
120601 TGCACTTATA GGAATACTCA CATGTATAGG AACGTTCATG ACAGCATTGT
120651 TCACAATAGT CCCAAACTGA AAATAACCCA AATGGCTATC AACAATGGGA
120701 TAGGTAGGTA AATTACAGTA TATTCATATA GCACTAAAAG TGAACAAACT
120751 TAACTACATG TAGCAACTTG GATAAATCTT ATACACATAC CATTGAGTAA
120801 GAAAAGTAAG ACACCAAAGA ATACAAGGAA TACGATTTGA TTTAATAGGA
120851 TTTAATTTAA TGGAATTTAA TAGAATACAA GGCATAGATT TTTTTTTGCT
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FIGURE 3MM

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120901 TTGTTAGTGT TTCCTTTATT ATAAAGCACT GAAATAAATA AATAGGTAGC
120951 TAGCCAATTT ATCCACAGTT TCTGGGAGCT ATATAAGATA GGCAAAGCTA
121001 AACTATTGTC TAAAAATATG TACATAGATA TTGATCTATA TAGAAAAACA
121051 AGAAAATTAT TAACATAAAA TTTAGCACAG TGACTTCTAG GGTTATGAAC
121101 AGAACAGGAC ACAGTGATGG GGACAAGATT CTATTTCTTG ACCTGTATCA
121151 TGTTTATGTG GACATTTGCT TATAACTGTT TGCTAATTCT GCAGTGTTTT
121201 ATTTACTTTT CTGAATATAT GTATAGAAAT ACATAATGAG CAATACCAAA
121251 CAAAATACTC AGTGGCTTTT TTGAAGGACA CTTAGCCCTT CTCTGACTCT
121301 CTTAGTACTC TCTTAGGTGC AGGGAATCTT CTGGAAGGGT TGGTGAAAGC
121351 CCTTCAATAT CTTCCTGCTC TGGTTTCTCA GCTATTTGAG GGCTCAAATA
121401 ATTACTCGTC TGTTATGTTT TTGTATGTTG TCATAAGGTT TCTTCTTAAT
121451 GTTCCACCAA AATGCTTCAG TGCCTTGCAT ACCATGAATA TTTTCTGAAT
121501 GAATAAATGT GTATTAAAAT GTTTTAATGC CTGAAAATAG ACCAGGTAGA
121551 AGAGGATGAA AAAGAATACT GGATAAATAA AGCTGGAAGA AAGAAAGAAA
121601 GTGAAAAGAA TACTCATGTA AACCCCAAGG ATAATCCAAT ATGACAGATA
121651 CATAACTTGT ATAGAGTAAT GTTTATTCTA TTAGGCATTT TCTTAGCACA
121701 GTGGCTCTGA TTATCCCTCA AAGTTCTTTG TAGCTTCTCT GAGTGACGTG
121751 TCTGTCACCC ATCACCTGGG GACTATCTGA TATGACTTGT TGTGAGATAC
121801 TGAGAAGGGA GAGCAGAAAT ATAGTCCATC CTGTCTGTGG GAGTAGTGTG
121851 GGGTCAGGGC CATTACCTCC CAAATTGCAC TGGGGGCTGT GACTTGCAGA
121901 AAGGATGCAG TGATTCATGA AAGGTGAATG CACTAGGGAA ATAGCCCTCC
121951 TTATTCCTGC TGCATCAAGC TCTTATAGTC AGGGCCAGTC CCGGGCATTG
122001 GGATGTAAAC ACTCTACCTC TCTAGTTGGA TGTTGTTCAC AGGATTTTAC
122051 TTAAAAAGAA CATGAGTGCA CTGGGTAGGG AAAACCTGTG TGTGCAGGAC
122101 CCATGTCATA CCAGTTTCCT TTGCCCAGAG CCAGCACTTT ATACAGGAGG
122151 CTTGGGATCA ACCATACAAA TCTTTCAACT AGGTCAATTA TTATGAATGT
122201 TTGCCTCTCT AGAAGCCTAC CCAATGTTTC TGAGCACTTT ATAAGTGCTA
122251 GGCACCATAC TGAGATTTTG ACATGGATTA TCACTGTTAA TTTCTAACTC
122301 TATAAAGATT GCCTTATTGG CTGGGTGCAG TGACTCACAC CTGTAATCCC
122351 AGTACTTTAG GAGGCCAAAG CAGGTGGATC ACCTAAGCCC AGGAGTTCAA
122401 GACCAGTCTG GGCAACATGG CAAGACCCTA TCTCTACAAA AAGCACAAAA
122451 ATTTTACCAA ATGTGGTGGT ACCCACCTGT AGTCCCAGCT ACTTGGGAGG
122501 CCAAGGTTGG AGGATCACTT GAGTCTGGGA GGTCGAGGCT GCAGTGAGCC
122551 ATGATTGTAT CACTGCAATC CAGCCTGGGC AATGGAGTGA GATTCTGTCT
122601 CAAAAAAAAA AAAAAAAGA AAAAAAAAG AAAGAAAGAA AGAAAGAAAA
122651 AAAAGGAAAA GAAAAGGGAA AGATTGCCTT ATTGTTCTGC TTTTGCTGTT
122701 TCTCAGGCTC TGCCAACTTG CTCAAGGTCA CAGTAAGTGG TGAAGGTAGA
122751 ATTTGAACCC AGAGAGCACA GCTCCAGAGC TAATGATCAC AACTATTGCT
122801 TGAGCAATTG ATTTGTTCAT TCATTCAACA AATTTCTCTC CAGTGATTCT
122851 GAATGCCAGA TTCTGTATTA GACAGTAGGA ATATGGTGGT GAGCATGCAG
122901 AAGCATTCCC TGCCTTTGCT TTGTGCTTCA TTCTCCCTAT TACATCCCTC
122951 AGGAGTTAGG TTTATTCTTA GAAGGGTAAG TAAAAGGTTC ATAGTGTGTC
123001 AAAGTGCTTA GAGAATGCAT AACTTGGGGT CCTCTCTGGG GGTAAAATTG
123051 ACTGTAGCTC TGCCTTCCAC TGGAATCAAT TGAAAGAACT ACAGTTACAA
123101 AGTGTAAAGA ACCCACAGCT GTTGTAAAAC CTTACACTCT CCAGAATGCT
123151 TGCTCCCTCT TTTCTCCCTC CCTCATCCCC AACAGATGGC TGCAAGTGCT
123201 TCCCTTGCTG CTTCCAGGTG ACTCTGAGAT AGAGAGATTA TCCAATGTAT
123251 GCTGTACCAA ATTCTGCACG TTGTCTGCGA CTGTTATAGA AATTTAGATC
123301 CTTTAGTTGA AACCTTCCCA ATCAAAACAA ATAACATCTT CTTAGCCTTC
123351 TTGATTTCAG GGTGAGCCAC ATATTTGAGG CCCAATAGGA CCCAAATTTT
123401 AATCGGTGCA TGATCTAAAT AAGCGAAGAG TTTATCCATG AAGGCCTATG
123451 CATGCCTGTG TGTGTTGACT GATGAATGAG GCTACTGAGA GAGATTAGAA
123501 AATTAGAAAT GTTTGCCTGC TGTGAGCAAT CTAGCAACGG ATGATAAACA
123551 TCCATAAAAG TGTTTATATT TTTGATCCTG GTAATTCTCC TTTGGAGGAA
123601 CATGTTGAGA AAATATAATA CTAATGTCTC AGGGAATCAA ACTGGTTTAA
123651 TTTTTCGTGT TTTTCAGCAC CTGAGATGTT CAGCTCCAGA AAAGGAGCAG
123701 GCTATTCCTT TGCTGTTGAC TGGTGGTCCC TGGGAGTGAC GGCATATGAA
123751 CTGCTGAGAG GCCGGGTACT GTAGTAGCAT TTCCTCTTTG GTTATTTTTC
123801 CAGCAAGTTC TATTTTAGAA TGAAAGAATG TATTGTTTGC TAAGATCCAA
123851 GCAGTTCACT TGAAAGCTGA AATCAGCTAT GCCATGTGAT GTTGATAACA
123901 CCCCTTGAGA TTTCTGCATA GGTTAATTCA TTTGTCCCGC ATATGGGACC
123951 AACCATGTCA ATTACCATTA AATTACACAG TTAAAAGTAA AGGAATAATA
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FIGURE 3NN

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124001  TGGATATTAT  AAACTCCCAA  AGAGGGGAAA  TCAATACACC  TCACTAAATA
124051  TCTTGTGTAA  ATATCTGTGT  TTGTTTAAAG  AAAGTCATTT  TGCAGTCATA
124101  GTACAGGACT  CTAATTCAGA  CATACCTCAC  CAAGGCTAGT  GTGAATTATT
124151  AATACAACAC  AATTCATGCT  CTGTCTTGTT  GGATTTCTAT  CACTTGGCTC
124201  CTGGGTTCTG  GGTTCAGTGA  CAAATTAGAG  TCATTTCCTT  TTAAAGGAAA
124251  CATTTCTTAA  ACTAAGAATC  TCTTTCCCAG  AAAAAAGAGA  TGAAAAGAAA
124301  GCAAATATGC  TGAAACATAT  TTTATACAAT  TTGTGCAAAC  TATTACATAA
124351  TAGAAATACA  CTCCTTAGGT  TATATCTCAG  TCAGCTCTGC  TTACCATAAT
124401  AAAATACTGC  AGACAGGATG  GCTTAAATAA  CAGACATCTA  TTTTCTTGGT
124451  TATGGAGGTT  GGAAGTCTGA  GATTAAGATG  CCAGAATGGT  TGGGTTATGG
124501  TGAAATCTCT  TTTTGGCTTG  CAGATAGCAG  CCTTTTTTCT  GTGTCCTCAC
124551  ATGGCAGAGA  GAGATCTTTG  TCTTCTTATA  AGTCTACTAA  TCCCATCACG
124601  AGGGACCTAC  CCCCATAAAC  TAACCTAACC  CTTATTCCCT  CTCAGAGGCT
124651  CCATTTCCAA  ATACCATCAA  ATTGAGGGTT  AAGGCTTCAA  CATCTGAATT
124701  TTGAGTGGGA  CACAAACATT  CAGTCCATGA  CATTCTATCC  TTGACCCCTC
124751  CAATATTCAT  GTCCTTCTCA  TATGCAAAAT  ACATACATTC  AACAGTCCCA
124801  AAAGTCTTAA  CTTATTCCCA  TATCAACTCT  AAAGTCTGAA  GTCCAAAATC
124851  TCATCTAAAC  ATCATAGAAA  TTGTGTATGG  GTGAGACTCG  AGGTATGATT
124901  CATCCTAAGG  CAAAATTTCT  CCTCAGCTAT  GTACCTATAA  AAGCAGACAA
124951  GTGGCCAGGC  ACTGGCTCAT  GCCTGTAATC  CCAACACTTT  AAGAGGTAGG
125001  AGGCAGGAGG  ATTCCTTGAG  CCCAGGAGTG  TGAGACCAGC  CTGGGCCACA
125051  TGGGAGACCC  TGTGTCTACA  ACACCTTTTT  TTTTTAATTA  GCCAGGCATG
125101  GTGGGGCAAG  CCAGTGGTCC  CAACTACTCA  GGTGGTTGAG  GTGGGAGAAT
125151  CACTTGAGCC  CAGGAGGTAG  AGGCTGTAGT  GAGCCAAGAT  CATGCCACTG
125201  CACTCCAGCC  TGAGCTACAG  AGTGAGACCC  CATCATTAAA  CAAAACAAAA
125251  CAAAAAACAA  ACAAACAAAA  AACAAGCAAG  TTATGTGCTT  CCAAAATACA
125301  ATGATACCAT  AGCTGTGGGA  TAGAGAATCC  CATTCCAACA  TTTCAAAAGA
125351  GAAATGGGAA  AGAAGGAAGG  GGCATCAGCT  CCTAAACAAG  TCCAGAACAT
125401  ATCAAAGCAA  ATTCTATTAT  ATCTTAAAAC  TCGAGAATAA  TCTTCTTTGA
125451  GTTGTTGGTT  TGCCCTCTAG  ATCTACACAG  GCATGGGAGC  AATCACTCTC
125501  ATGGCTGGGG  ATGGGGAGAG  GGGACTTGCT  TAAGTGGCTC  TCTACAAAGG
125551  CACTACCCAC  ATGGCTCTCT  GTGAAGGCTC  TGTCTACACA  GCTCTGTTGA
125601  GTGGTGGTCC  TGCCCTTCGA  AACAGAGGTG  GAGGCAACCC  TGCTCCCCAA
125651  GCCAGTGCAC  TCTGGACCTG  TAGTGGGAAT  GGCAGCCCTG  ATGATCTGTG
125701  AATCGCCCTC  ATGATCCTTC  TTCCTTTTAC  TTGAAGGATA  GCACATGTTC
125751  ACAGCTGGAT  AGCATTACGG  TCCCAGCCTG  TAAAATCCAA  GAAGTCTGAC
125801  AGCCTTTCTT  CATAAATTCA  AACTGGCAGC  ATCTGCTAGT  ATAATCCCAT
125851  CTTTATTTCT  AGCTTCTGTT  GTGATAACTA  CTTGATTGTT  CAGCTACACT
125901  CTAGTGTGCT  CTTCAGAACA  GGCTTGCTCA  TTTTCTGCAA  TATGGATAGA
125951  AATCTTCAAT  TTCTGGTTGC  TTTTTGCTTA  ATTATTTTTT  CTTCAATTCA
126001  AACATTCCCT  TTAACATTTT  ACTATAAGCA  GACAGAAGGA  ACCAAGTTAC
126051  TCCTTCAAAG  TTTTGCTTAG  AAATCTCCTC  GGCTGGCCTG  GTGCAGTGGC
126101  TCATGCCTAT  AATCCCAGCA  CTTTAGAAGG  CTGAGGCGGG  CAGATCACCT
126151  GAGGTCAGTA  ATTCGAGTCC  AACCTGATCA  ACATGGAGAA  ACCCCATCTG
126201  TACTAAAAAT  ACAAAATTAG  CCGGGCATGG  TGGTGGATGC  CTGTAATCCC
126251  AGCTACTCAG  GAGGCTGAGG  CAGGAGAATC  ACTTGAACCT  GGGAGGTAGA
126301  TGTTGCAGTG  AGCTGAGAAC  ACAACATTGT  ACTCCAGCCT  GGGCAATGAG
126351  AGCGAAACTC  CATCTCAAAA  AAAAAAAAAA  AAAAAAGAAA  TCTCCTCAGC
126401  TAAATATCTC  ATTTCATCAC  TCACAATTTC  TACCTTCTGC  AAAATAGTAG
126451  AACACAGTTC  AGACAAGCTC  CTTGCCACTT  TATAACAAGA  ATCACCTTTC
126501  CTCCAGTTTC  CAATAACATG  TTCCTCATTT  CTGTCAGACC  TCACCAGAAT
126551  CACCCTTAAT  ATCCATATTT  CTAGTGCATA  CATCCACAGT  CTTCCAGCTC
126601  AATAACTAGT  TCCAAAGTCA  CTTCCACATT  TTAAGGCATT  TGTTCCAGCA
126651  GCATTCCAAT  TCTCAATACC  AAAATTTTAG  TCTGCAATAT  CTGCCTTCAC
126701  AAAATACCAC  AGAATTGGTG  GCTTAGGCAA  CAGAAATTTA  TTTTCTCAGT
126751  TATGGAGTCT  AGAATTCTGA  GATTAACGTG  CCATCATGGT  TGGGTTCTGG
126801  TGAGGGCTTT  CTTCCTGACT  TGCAGACAGC  TTCTTTCTTG  CCCTCACATG
126851  ACGGAGAGAG  AGATAATCTC  TTTCTCTTCT  TTTTGTAATA  AGGCCACTAA
126901  TCCTATCCTG  AGGGCTCCAC  CCTTATGACC  TAATCTAACC  CTAATTACCT
126951  CCCAAGGGCT  TCATCTCCAA  ATACCATCAT  ATTGAAGGTT  AGGGATTAAA
127001  TTTAGAAATT  TTGGGGGGAT  ACATTCAGTC  TGTAACAGGT  TGTATACTCT
127051  CAAGGTCCCA  GTGATGGATG  CAATCAGTGA  TTCCTCTAAG  ACCAAAGAGT
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FIGURE 300

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127101  TGAAGACCTG  ACTTTAGGAG  CTTGTTTATC  CCACAGAACT  AAAGAATTGG
127151  GTATCTCAAG  TCATCATCCA  GATACTGCAG  CTCTCCTCTC  CTAACTTTTT
127201  GGAGTCATTC  TTTCTGCTGC  TGTCAATAGC  CCTCTTCTTT  GGTCCCACAA
127251  CACACCATCA  TGATTTCTGC  ATTAAAAATG  CCATCTCCCA  AGTAATTAAC
127301  CTATTCACAG  TAAGAACAGT  TGTTAGAAGT  TGGGGTTATT  TCATCATGGT
127351  CCAATGGCTT  TATCTTGCTC  AGGAAATCAA  AGATGAGTGT  TTCTAAAGCA
127401  AAAAAAAGGA  GGATCTCACA  ATTGTATCTG  TTTCATTCAC  TCTGCAGGGT
127451  CCATTTTACA  CCCAAACATT  CATTAGTTCA  TTGTTTGTAC  TCCTGCCTTT
127501  CCTGAGGAAG  TCATTGTAGC  ACTATTTCTT  AAGTATATTC  AAATTTGGAT
127551  AAGTTAGTCA  AATTGATGTG  AAAGGACCAC  CCTTGTAAGC  CAAATGTGTA
127601  AGTCCTACAT  AGGGATATTA  CCTGTTTTTA  TCTCCTGATG  GGCTTTTTTT
127651  TTTTCAAGTT  TCTAAATAAA  TCCAGTGAAC  AAGTAGATAC  GCTACTCATG
127701  ATTATATAGG  AAAACAGAGA  AGAGAAACAT  ACACTTACTT  AAAAGTAGAA
127751  ACATATCTGC  TCTTTCCCAC  TTCACCCTTA  ATTTTTTTCT  CCCCAGCCAA
127801  TTTACTCACC  TTCTGTGGCT  GTGCTTCTGT  GTTAGACCCT  TGCTAGCTGC
127851  TTCTGGGGTT  CAGAGCAATT  GTGCTCTGCC  CTCATCTTTT  ATGACACACC
127901  TAGCAAAACA  GAAGCAGAGG  AGCGAGTTGA  AACAGACAAA  CGACTATCTG
127951  TTATTCTTCA  AACATGCCTA  GGATTGTATT  TAACTATCAC  CTATCTAAAA
128001  GAGGTATTCT  CGCCTGCCTG  GAAAGAATTT  TGCTAAGAAA  ATTGTTTCTC
128051  TTCTTCCCAT  ATTATTTTAC  CTCTATGCTA  GTTCCCTGTG  ATTTGATATG
128101  TCAACTTTGA  CAAATTCATT  TTTCTAAAGC  ACAGATATGA  CCTTTTTTGT
128151  TAAGAAAAAG  AAACTACTGT  TGCTCCCCAG  TGCTACACAC  ACACACACAC
128201  ACACATACAC  ATACACACCC  TTCACAAGCC  TTATCTGCAC  CCCCGCCCAC
128251  TCCCCACAAC  AAACTTCAGA  TGTCTTAGCT  TGGCATTCTT  CGGAATTAGG
128301  TCAACGTTTC  AGATTTTGCT  TCCATTTGTG  TATTTCTGAC  CCTTCATGAA
128351  CTCATTTTGG  CCTCTTAGAA  CTTCTTCCTC  TTCTCAAAGC  ATCTCTTGGG
128401  TTTTTTAACC  TCTTGTTCCT  TCGCCTATAA  AGAGAGTTTC  CAAGGCAAAC
128451  CTTGGTCTTC  TTTAAAAATC  ACTCTGCGTA  AGATTTGAAA  TCACTAAATG
128501  AAGTTTTAAT  AAAGGATATA  TCTTCATTGC  AGGGCTTTTC  AAAATCTTTA
128551  TAGCCAAGTA  TTTTGGTCAT  TTCTAAGAAA  GGACACACTA  TTAAACTATT
128601  CCAGTTCGTG  TTGGGGAGGT  TTTTCTAGAT  CTCTTTATAT  TCAAATTCTA
128651  TTCATACTTT  ATCACCTATG  ACAAAATAGC  ACTTTCTCTA  AAGAAACATT
128701  CTCTGACCTC  CCTATCTAAA  GTGATCCGAA  TCTCTTCCAA  ACATTTATTT
128751  ACTTTATGTA  TCCTGTGAAT  CTTTGGAATC  TAAGCTTATT  AGAAAATATA
128801  GAAAACCACG  AAAATGAAAG  CAAAAATCAG  CTGTAGTCTC  TAAGGCAAAG
128851  AACATTTCCA  ATTAAGAAAT  TAAACTCCCT  TTGACTTTTA  AACCCCATCT
128901  TAGCAGTTTG  TTGCATTCAC  TTCCAACTTG  TTTCTGTTCT  CATAAGGATA
128951  CTCTATCTTC  AGATAGATAG  ATATAGATAG  ATGTGTTGTT  TTAGCAAAAA
129001  TAGAAGTATG  TTTTACCTTG  TTGAGCCTTT  TTTTTTTTCA  TTTCATAAGA
129051  TAAAATGTAC  AGCTTTCTAG  ATCAGAACAC  CTAAATCTAT  TTTCTTTTTA
129101  AGGATTAAAT  CTATAGGCAT  ATCAATTTTT  ATTTTTTATC  TCTTGTATAT
129151  TATTAGGTTG  TTAATTCATT  AAAGGTAAAG  TATGTATCTT  ATATAGGTTA
129201  GTATTATTCA  CAGTATTTAA  CTGTTTTTTT  TTTCCTCAGG  AGAGTCTTGC
129251  TCTGTCCCCC  AGGCTGGAGT  GCAATGGCCC  AATCTCGGCT  CACTGCAACC
129301  ACCCCCTCCT  CTGTCCAATC  AACCCTCCCG  CCTGAGCCTC  CCAAGTAGCT
129351  GGGACTACAG  GCATATGCCA  CCATGCCTGG  AAATTTTTTG  TATTTTTTGT
129401  AGAGTTGGGG  TCTTACCATG  TTGCCCAGGC  TAGTCTTGAA  CTCCTGGGCT
129451  CAAGCAATCC  ACCTGCCTTG  GCCCTGCAAA  GTGGTGGGAT  TACAGGTGTG
129501  AGCCGCCGCA  CCTGGTCACA  ATATTTAACT  TTAAATAGGT  ATATAATACA
129551  TGGTTATTTT  CACTCACATC  CATGTGAAGA  GACCACCAAA  CAGGCTTTGT
129601  GTGAGCAACA  AGGCTATTTC  ACCTGGGTTT  CAGGTGGGCT  GAGTCCGAAA
129651  AGAGAATCAG  CGAAGGGAGA  TAGGAGTGGG  GCCGTTTTAT  AAGATTTGGG
129701  TAGGTAAAGG  AAAAAGGGGG  GTTGTTCTCT  GGTGGGCAGG  GGTGAGGATC
129751  ACAAGGTGCT  CAGCGGGGGA  CGTTTTGAGC  CAGGATGAGC  CAGGAGAAGG
129801  AATTTCACAA  GGTAGTGTCA  TCAGTTAAGG  CAGGAACCGG  CCATTTTCAC
129851  TTCTTTTGTG  GTGGAATATC  ATCAGTTAAG  GCAGGAACCA  GCCATCTGGA
129901  TGTGTATGTG  CAGGTCACAG  GGGATATGAT  GGCTTAGCTT  GGGCTCAGAG
129951  GCCTGACAGT  TATTGAATGA  ATGGAGAAAC  AAATCACTTA  GACACCTTCT
130001  AGGAAAAAAT  GACCAACTAT  GCTACCTGCA  ATTACGTTTC  AAAATGTAGC
130051  TTATCTGAAG  AAAAGGAAGT  AACATTTAAT  TACAAGCATC  AATACAACTC
130101  AAGCACAGAG  GAAGTGTGCT  AAACAATTTC  CTCCATACGT  ACAAATTTTT
130151  ATTTACAGAA  AAGTATATGT  CTTAATGAGA  AAATGTGCTC  GAAAACATTC
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FIGURE 3PP

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130201 TCATCATTTC TGAGTTTGGT TTCAGTCTTA ATGAATGTGT CCCTTAACTA
130251 TTAATCTGCT TTGTCATCTC TCTAACTCCC TACTATCTCA TTGCCATTGC
130301 AAAGGCAAAG GTCCACATCT TTTATAGTTT CATATTATCC AAAAGTGTTA
130351 ACTTAGGATA GATGTGTACA TAGTTTTGTA CTCATTGTAC ATGCTTAGCT
130401 GCAATTCTTT TGCCTTTGCA CTTCTGAAAT ACAACCATAT TCACAACACA
130451 TCATTTGTTC CCTTATAACA TTTCACCTTT TCCACTTTGT TTATTCTCTA
130501 TATGCTCACT GTTAGTTTAG ATGCTGCCTT AGGCTTTTAT GATATATACT
130551 GTGACTGCAT ACTGTAATTT TTCTCTATAG CATGTATCCC ATTTATTTAA
130601 GTGTGTGTGT GTGTGTGTAT ACAGTCTATA TAATAAATTT ACATGCTTCC
130651 TTAAGTAGAC TGTAGGCCCC ACCAACATAG AAACCATATG TGTCTTGTTC
130701 TTCATTGTAC CCTCAATGCC TAAGAAAGGT GCTGGAACAT GGTAGGCATT
130751 CAATAAATAA TTGGTAAATA AATAAATATA CAATTCTGGT AGTTGATTAA
130801 TTCAAATTAA TTTTAAAATT TAGAACTGTA AAAGTAAATT AAAAAAATAAG
130851 ATAAAGACAA TGTGATTATT TTTTAATAAA CCAACAGGTC ATGGAGATTT
130901 TAAAAATTAA ATTCAGTCAT ATGGCCTTGT AAAGTAACTA GAGAAAAATG
130951 TACACACTTA AACCAGCTGC TTGTGGCATT CATCAGTTAA TTCATTTGTT
131001 TATAAAATCA TTTTATTTTC TAGGTGGCCC AGAAACAGTA GGTTGAGAAG
131051 CAGCAATGAA TTAAAATCAA GAAGAAACAC AGAAAAAAGT AAAAACACAT
131101 GTGCATACAC ATATAAGCCT AGAAGCTTGA GTATACTAAG CCTAATCTGA
131151 TTCTTAATGA TAAACATGGT CTGAATCATA TGGAGTAACC TAACCCTTTG
131201 GCTACTAAAT TACCAATAAA CATTGATAAT GGTGATAAAG CATCTAGCAC
131251 TCCTTTACTG ATATTGAGTT AATGAGTTAT TTCTACTATA TAATTACCAA
131301 GACATATGAT ATAGCTATGG TCCTTTATTT AGTGTTGAGG GGGTAAATAT
131351 GGCAGTTGTT TTTAGATCTT ACTTAAAAAG CAAAAATGTT TGAATTAATC
131401 TCCCTTTCAA GGGCCACCTC CTGGCACTTC ATGGTTCCCAT GAATAGCTGA
131451 CATTGACTTG CCATGTGTAA AATTAAGCTT TTCTTCCCAT CACTTTTCTT
131501 GAGGACTCAT TTTGCTGTTC ACTATTCATT CACATTTACA TATGCCCATT
131551 TTTACCTTTG TGTCAATAAT GATAAAAATC TCTCTCTTAT ATTGTGTCTA
131601 ATACTATTAG CCACTCACTC TGTTGAGAAA TTTACACATA TTATCTCCTT
131651 TAATTTTTCC AGCAATCTCA TGAGGTAGCT CATTTTACAG ATGAAGTAAC
131701 AAGCTCAGAA ATTGAGTGGA GAAGTTTAGC ACCAAATCCT TTTAACCTCA
131751 AACACATGAT TATTTTATAT TACCTCTTAA CACTGATTTA CTACAGGGAA
131801 AAACTTAAAC CCTTTCATTT CCCCCAATTT AGGTCATCCA TCAACAGTCA
131851 TTTATTAAAT ATCTTAAAAG GGCCAGGCAT GTGATCAATG TGTATATCCA
131901 TATTAACTGT GCTGTGGCTA GTTAATCGAA TATGGAAATT TTGTTCATTA
131951 AATAAACATG TATTGTGCAC CTACTGAATG CTTGGTCTCA TGAACAAGAA
132001 TGATATAATC TCTGGCTGTG AGTATCTTAC AGTTCACATA AGAGACATGA
132051 AATTTCAGTG TTGGTGAGTC CCCTACAAAA TAATATAGAT AAAGGCTGTC
132101 CTCTAGTGTA AAGCTGTGAA AACTACAGCT AATCCACAGT TTTCTTTTGT
132151 TTAATTTCTT TTCTTTTTTAA ATTACTTTTC TTCAAAATTA AAACTGTAGA
132201 AGAACCTGGT TCTTCCCCCA AAATTTTTTT TAAAAGCTTC TGCCTCATCA
132251 CAAAATTCTC CACCCTGCCA TACTCTGTGG AACCAGGGAC TCATAGCATT
132301 TGTGGGACTG GAGTTGATGT TTTCTGAGCA GTTTTCTGTC CTGAGCTTCC
132351 TCATTATGTT GCAGTGAAAG GGATGGTATG GTAAAATTCT GGATTTACTT
132401 GCAATCAACC CTTACATAAT AATTTTTTAG ACTTCCATTT ATTGAGGACT
132451 TGTCCAGTAT TTCGTGTTAA TACTTATATA ATACCTTATA AAACAATTTC
132501 AAATCAGCAT CTCAGAGGCT GATTCAGTCC ACTTGAATGT TTTGTTTGGC
132551 TCAGTGGAGT GTTCAACTTT AAAATTTATG GTATTTTAGA AGCGACCATA
132601 AATTCCTAGT GTCTCTTTAA GAAAAAGTAG GGGGTCTGGC AACACAGGAC
132651 CACCTACACA TATGGCAACG CAAGAGTCAG CTGGACAGGG TTAGAAATTG
132701 ATATAGATAT TTTATCGGTT GAAAGTTTAG CTTGGAAACA TTTGGAAATT
132751 TTTTTTTTCT TTTGTCCTAT ACAAATGAAG ACTTTTACTT CTTTTCTCCC
132801 TTAAGAGACC GTATCATATT CGCTCCAGTA CTTCCAGCAA GGAAATTGTA
132851 CACACGTTTG AGACGACTGT TGTAACTTAC CCTTCTGCCT GGTCACAGGA
132901 AATGGTGTCA CTTCTTAAAA AGGTAAGAAG GAAGACTGCA TGTCCAAACG
132951 AAGTAACAAA AGGAAGCAGG CTCTCTGGCT TAAGTTTAGA AGTTAGTATA
133001 CAATATTGGG GACAGTCATG ATAGTATACA TTTGTAGAGT GTATTTTCTA
133051 GCTGTTAGCT TTCAAATACA TGGCTTCATT AACTCAACTC AGATTCCCCT
133101 TGGATGTCCC AAAGCCATCT TAAACTCAAA GGACTTCTTT ATGCTTTGTC
133151 TTTCCTGAAT ATCTTCTCAG GAAATTACTC TCAGTGACTG GCTTCTCTAT
133201 CCAAATCCAC TTACGCCAGC CAGCAACCAG GACTCATCTT GTCATACTGC
133251 GTATTCAATT NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN
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FIGURE 3QQ

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133301 NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN
133351 NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN
133401 NNNNNNNNNN NNNNNNNNNN NNNNNCTAGC CTTATAACGG GTTTGTCCAC
133451 ATACACTTTT ACCACTCCAT TCTATTCCCC ATGCAGCCCC ACAGTGGTCT
133501 GTTAAAGGAC AGTCCAGGAT ATTTTCCTTA TTCTTAGAAT AAAGATTAAA
133551 ATAATTTTGT GGTACAAAAG TTCAAAATAC CTCTCAAGCC TTGTTTTGGA
133601 CTTTTGGACT TTTGTCCCCC CTTTGACTAC ACATAAACTG CTTTGGCCTT
133651 TTTCTTCTTC TTTTCTTTCT TTTCTCCTTC TTCACTTTTA CATACCAGTC
133701 TTCCTCTCAC CACAGGACCT TTGCACATGC CAGTACCTAT TCCTGGAACA
133751 GTGCCTCCAA TCCTAGTTCC TCCAGTTCCT CCTTGAGAGC AGTACTACTC
133801 AATGTGGTTC ACTGGTTCTA GTCCATGAAT TTTTTCTGCA GGTCTATTGT
133851 AAGTAAAGAA CTTGAGAGAA GCATTTAGAA ACTTTTATAG CAATTGGACA
133901 CTGCTGTAGC ATCTAAACAC ATGATCAATG GACTTATCTT ATTGAAGAGG
133951 GTCCAAGCTT GTTTGACGGT TGTTGAACTC AAGTCACAAG GTGTCTATGT
134001 GGGGTGCTGC ATACTGGCAA TGCATAATAA GACCACATAC TGATTTCAGT
134051 GGATTGGAAA TTGAAACAGT ACAAAAACAA ACAAAAATAA CTGACCCTTC
134101 TACATAGTTT GGGAAGCACA ACTTTAGCTC TTAGCTCAAA TATCACCTTC
134151 TTGGTGTAAG TTCACATAAC ACTATCTTTC CTTCATAGCA TTTTTCAGTT
134201 TAAAATTATA CCCAGCATTT GTGTGATCCT TGGTTACGTA CCATTTTCTT
134251 CTTAGCTTCA TGAGGGTAGG GACCATGTCT GACATGTGTT ACCATTGTAT
134301 TCTCAGCATC TAACACAAAG CCTGAGAAGT GAAATTTGAC AAGTATTCAA
134351 ATAAATGAGG TCCACAGCTT TCATCAGATT TTCAAGGTAC CCATCTTCAT
134401 CAAACAGATG AAGAACAGTT ATAGCGGGAG GTCAAAAGTG TATATTGAGT
134451 GATGATACAA AACAAGAATG AGGGGCCCAA GAGGAATGGG CTTGGCCTTT
134501 TTTTTTTTTT TTTTTTTTTT TTTTTTTGAG GAGAAAATTG CACCAGTTGT
134551 GGCTGGTAAT GGAAAATAGC TTTAGTGGCT AAGGAGTCAT CATTTGTGTC
134601 TCTTGTTTTT GGAGTCAAGT TCCTTATTTT GGAATAGGGA CATTGCATCA
134651 GTAATGTCAA AGACATAGAA TGGGGGATCA TTTTTCATAA GCAAATTCTG
134701 CTTAGTTCCA AGACAGCCCT GCTTCACTCC ACAAATTACA CCCTGAGGTT
134751 GCATGGTTGT CATCTTCAGA AGCATTCTCA AGTGGGACTG ACAATGCCTA
134801 TTTGAGCCAC ACAATTGCTG TGATGTTGGC TCAGGAATGG TTAAGGGGGC
134851 AAAAATCTTT TATCTCAATT AGTAAAATCT AGAACTATAA CAGTTACTTT
134901 AGTTACACCT TATCTATGCC GCCCCCAATG TATTTTAATT AGTTGTAAAA
134951 ACAGCTACAA TTCTTAGTAG GAAATGAGTT CTACTTGTGA AATGTATCAA
135001 CATTTGTCAC CATAGGTTTT CTACTAGGTA CTTTGTATAA ATAGCCTCCC
135051 ACTAATCCTG ATTACAATCG TATGAAATAC ATTATTACCA CTTTTTTTAA
135101 ACACATGGGT AAACTANNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN
135151 NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN
135201 NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN
135251 NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN
135301 NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN
135351 NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN
135401 NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN
135451 NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN
135501 NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN
135551 NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN CATTCCAGGT TTATTCCTAA
135601 TGTGAGTCAA TCCTAACAAA GCCAAATAAC TCCCATTCAG TGCGCATTAC
135651 CCGGTGTGCC AGATTCACAC ATTGTCTCAT TAGATAATCA CACCAGTGTT
135701 GTTAAGAGAG CCCCAATTCC CATTTTACAG TATGAGAGAA TTGAGACATG
135751 CACAGGGTAA GTTTGTCACC AAAGGTCACA AAGCTAGCAA GTAGTCAAGC
135801 TGGGATTCTA ATCCAGGTGT ATTTGCGACT GAAGTTCTAG CTTTTAACCA
135851 CTTTTTATGG TCTGTTTTTA TTGAAAGGAA GTCCTAGTTC CCCAAATAGT
135901 CATTCTCATG AATCTGCTGG GGTTTTTTTT AAGTTTTCTT TGATTCTAAA
135951 GATGCAGAAG TTTGTGTCCC TAGAGATCTG AGTCAAAGAA TTGAAAATTG
136001 TTGGAGTTGG GGTGAGGAAT TTATTTTAGC ATTTGCCCCT CATCCTTTGT
136051 TTGTTCTGTC TCAGGGATTT ATATTTGTAA GGACTGATAA CCAAAGACAT
136101 ATAATTCCCA TTGGATGGAT AGCCAAACCA ATGGACTTCT GTGGTCTACT
136151 GCATTATGCT GGTAAGAGCC AGAGTCCAGA AGCTTAGGCC AAAGGTCCCA
136201 AGTGAGGCCA CTAGCTCCTT CTCTCTGCCT AGAACTGAAA TTATATGTTC
136251 AGTTGTAGGT ATATTGGGCA GAATAAGAGG CTTCTAAAGG GGCCTGTAGA
136301 ACCAATTCAG TTTTCTGTTT TGGCTGTCAT GGCAGCTCAG GCCTGCAATC
136351 TCAGCACTTT AGGAGGCCGA GGCAGGAGGA TCAGGGGTTC AAGATCAGCC
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FIGURE 3RR

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136401 TTGGCAACAT GGCAAGACCG TGTCTCTACA GAAAAAGAAA AAAAAATTAG
136451 GCAGGCGTGG TGGTACTTGG GTGTAGTCTC AGCTACCTAG GAGGCTGAGG
136501 TAGAAAGATC ACTTAAGCCC AGGAGTTTGA GGCTGCATGA GCAGTGATTG
136551 TGCCACTGCA CTCTAGCCTG GGTAACAGAG TGAGACCCTG TCTCAAAAAA
136601 AAAAAAAAAT TACTCTTAAG CCCATATGAG GCATTTGCTG TGGGAATGTG
136651 AGAGTGTGAT CCTTCATGTA CACACAGCAG GAGGCATGCT CCAATGAGAG
136701 GGTAAGGAGA AAGTACAAAG TGAGAGAAAG GAGAAAGCAG GGTGGTGGAA
136751 TTGTACCTTA TGGAGCAACA GGAGGGTAGG TCTGAGTTCT TACCTCTCCG
136801 CTTTGTGGGG TCCATTAGGG GCAACTTGTA CCATAATTGA CACATGACAC
136851 AATGAAGGTC TAGGCACCCC AACTCTTGCT TCCCCCTCCT TCTATGTGTT
136901 GCGTCCCTGC AATTAGCCAT CAATGCTGGC TCAAAGAAG TTCTACGTTA
136951 TGCTTCTCTG ACTTTAGTGT GAATCGGAAT CATCTGGGAA GCTCATTAAA
137001 GTGCAAGTTC TTGGACCTCA CATTCTGAAA TTCTGATTTG GGAAGTCTGG
137051 TTGGAGAACT GGGAAGCTGA GCAAGCAACT TAGGTGATTC TGAGTTACAT
137101 GATTATTAGA GCGCACTTTC GGAAACATAA CCCAAAATTT ATTTTCCACT
137151 TTAGAAAAAT AACTGTAAGT CGGCTTTTGT TTTTACTCAT TGAGGCCTAA
137201 TTGAGAGTTT AGAAAAATAA ACGAAGAATA TGAAAAACGA TGCTGGCAAT
137251 AAATAACGTA AAACTTAGAG TGGGAATCCC AGTGTATTAT TCATGGACTG
137301 CTCCGTTAAG ACTAAGTATT ATTTTCCGTA TTAGGTCTGC TGTGTTTTTC
137351 AGAATGATAC AGTAATCTGA GGATTGAGCC AACTGTCTTC CTTGCAGAAA
137401 GGCAGGCTGA ATTGTGATCC TACCTTTGAA CTTGAGGAAA TGATTTTGGA
137451 GTCCAAACCT CTACATAAGA AAAAAAAGCG TCTGGCAAAG AAGGAGAAGG
137501 ATATGAGGAA ATGCGATTCT TCTCAGGTAA GCAGGTCCCC ACCAAACTCA
137551 GGGTCATGGG TATCCCCATG ATGGCTGCAA TATCTTCGAG AGCTTCTACT
137601 GGGAGGTCAT TTCAGCTTCC TGCTTTTGCT GCTTAGTGAA ATAGGAGAAG
137651 TAGATCAGCC GGGTTTCTAA AAGGGCAGAC CAGAGCTCCT CTGAGGATCC
137701 TAGCAGCAAC ATTTTACTTG TAGGCTTTCC GTCTAGAGTT CTGCCATTAA
137751 CTTGACTCAG TTATTTCTCT CTTCCAGTTC TCAATTCAAA ATTTACAAAT
137801 TTCCTGGGAG AGGAACTGTC ATTGGCCAAG CTTAGGTCAG GGGATGATTC
137851 ATAAAATTAT GGTAAAGGGG CAGGTTTCAA AGTACACACA TGGTTGTTTT
137901 GGACCTCACT CCTGCTTTGA GGAGTTTCTG GGAGCAGCCA ACCCTAGAGA
137951 TGATGTCTGT TCTTTGCCAC AAGCAGAATT TTATGATATC AAGCCTCACA
138001 GAAGAGTGTC TGTTCACAGG AATGACGGAA TTCTAACATG GTGGAGCACT
138051 ATTGCTGGAT TTCAGGCTGA GTTAAATTAA CTTTGTAACT AAGTATATTA
138101 TTCTCTGTCA GAGTCAGAGC TCAGATTTCA GTGAAGTAAC TTGCAAACAC
138151 TCAGTAGGAT TTTATACTCA CATGTGGCTC TATGAATTAT AATGATGATG
138201 AAGTAATAAA GTTACTTTGC CTCTAAAGGT CATCTATCTA TCCACACGAC
138251 CATTTCCATT CCTCCATCAA TCCCTGCCTC CCTCCATCCA TTCATTTAGG
138301 CTACTTTTTT TTAGTAGCTA TGATCTGCCA GGTCCTGTGC TAAAGACTGG
138351 AGTGAGAAAT GATTGAGATA TAATTTCTAT ACTCAGTGCT GTCCCTTTTC
138401 TCAAAGATTG TGTAGTCTTG TGGTAAAGAT GGCTCTGCAA ACAAATAAGT
138451 ATCCTCCATC TCCTTAATTT CTCTAGTAGT CAGGGGCCAC TATATATTTC
138501 AATGGACAAT TAACCAACGT TCACATCTCT GTCCTGTTTG ATCACAGAAC
138551 TGGCTTCTCG TCAGATTCCC TTCAGGAAAT ATTTTCTAGG ACCCTCCAAG
138601 GAATGCTTAG CTGTGCTGCT AACCCGTCTT GCATATTGCT TGTCTCTGAA
138651 CTGTCTTCTT CCCAATGGTC TGTTCCTCAT GATCATGTCA TAACCAACCC
138701 GCTTCTCCAG ACTTGCTCCT TCCCCTGACC TAGCAGAACT TGGCTCAAGG
138751 TGGATACAGG CCTCTCTGAT AACAGGACCT AACATGTGAT AAAAACCAAG
138801 AGATCCTTTT TATTACAAGT TTTTAAAGTT TTAGAAATAA CTGAGCAATT
138851 TAGGAATAAC TTTTGACCAT ACGTACCATG CTCAACATGA TCTGCCCATC
138901 TTTCCTGCCA CATCCTTGTA CTATCCCACT CTGACCCTCA CTTAAAACCC
138951 TCCAACCTCA CAGGCCCTGC AAGTGTCTCA CTCTCAAGCA CTGAACCTTT
139001 TGTTCTTCTT CAAGGCCTTT GCCCTTGCTC TTCCCTGTTC CTAGAATGGT
139051 CTTCCCTTTC ATCTTCACAT AGGGGGCTTC CTCTCATTCT TTATACCTTA
139101 AATATCACCT TGTCATTTCT GTTGTTGAAT TATAGGATGT TTTTTACATA
139151 TTCTGGATAT TGGACCCTTA TCAAATATGT GAACTGCAAA TAGTTTCTCC
139201 CTTAGTCATT CTACGAAGCC AGCATTACCC TGATACCAAA CTGGACAAAG
139251 ACATCACAAA AAATGATAAT TACAAACTGA CATCTGTTAT GAATATAGAT
139301 GCAAAAATCC TTAACATATT AGCAAGGTGT TCAGTTAGGC TTTTGACTTA
139351 AGATGTTTCT TCTTTTTTAA TATTGGTGTT TATAGCTATA AAGTTCCTTC
139401 TGAGCACTGC CTTCACCTAT CCCATAAGTT TTGGGATGCT GTGGTTTGTT
139451 TTTAATTCAT CTCTAAGTAT ATTCTGATAT CTCATGTGAT TTCTCTTTTT
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FIGURE 3SS

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139501 GACTCTTTTT TTAAGAGTTT GTTGTTTAAT TTCCACATTT TTGTGAATTT
139551 TCCAGTTTTC CTTCTGTTAT TGATTCCTAC CTTCATTCCA ATTATTTCAG
139601 TCTTTTTAAA TTTTTTGATA CCTGTTTTGT GGTTTCCTTC CATGGTTTCC
139651 TTTAACTCTG AGCATATTCA AGACGGTTGT TTTAAAATCT CACTCTAGAA
139701 AGCTCAATGT TTGAGCTTCC TCAGGACAAT TTCTATCCGT TGATTTTAAG
139751 TCTTTGAATG GCAATATTTT CCTGTTTCTT TGTGTGCCTT GTGATTTTTT
139801 TTCTGTTGCT ATTGAAAACT CGACATTTAA TATGATAAT GTGGTAACTC
139851 TGGAAATCAG GTTCCTCCTT TCTTCATGGT TTGCTATTTT TTGATTGTTG
139901 AAGGCTGTAG TTATCCATTG TTTAGCGACT TCTCCAAACA ATGTTTGCAG
139951 AGATTGTCTG CTTTGTTGTG TCATCACTGA AGTTTCTGTT ACTTTAGCCT
140001 GTGCTCAGCT AATGTTTTGA CTGAGATTTA ACACCAAGAG CATTTTTAAG
140051 TTGTTTTTCT TTTCTTAATT TAGTGTTCAC TTGGTTCCAG TAAACCTTTG
140101 AGTGCTTTCC GGAGTTTTGA CAAAGTTGGT TTTGACAGTA TCTGCTTGTT
140151 TTTTTGATGT TTCTGTTCAG AGATGGGGCT TGGAACTGCT TACATCAGCA
140201 TTTTTCTCTA GATTCTTCTA ATCTTGTACC CCAGGTTCAA AAATAAAAGG
140251 TACTTTGCTT CAAAACAAAG AATAGTCTTT CTTCCAAGAA GAATCAGAAA
140301 GATTATGAAC TATTTTTCTG ATTCTTCACT CTATTTTCTC TCTTTTACAT
140351 TAAGGCTTTT AAAACATGAG TCAATCTTAC CTTATTATAT TATTAACATG
140401 CTCGTTCATT CATTCATTCA TTTATTCAGA TGACTGTAAA ATTCCTGCTT
140451 TGTTAGGAAA TATTTCTGAC TAGGTGGTTA ATGCTATGGT TAGATACACA
140501 AAGTGCTGTG GGAATTGCTC ACTGGACCTG AGTGAAGGGG TAGGATAGGC
140551 TTTCCAGAGG AGGCAACATT TGATCTGGTT CCTCCAGATT GAGCAGAGGT
140601 AGGTGAGCAT ACAGGAAAGG ACAAGAGCAT TTCAAGGCTG GCACATCTCA
140651 GGGCACAGGC AGATCTTAAT GTTACAGAGG AAATAAAATG ACAGGTGGTT
140701 TCTGATCATA GGAATTACCC ATGCTGTGTT CAAAAGGCTT GTGACATTAC
140751 TCATCCTCCC TGCCTTTAGT CTTATCTAGA GCCATTCACT GAAGGCATTC
140801 CTTCAGCAAA ATCTAACAAG AACATACACC ATATCAGTAT CATATTAGCT
140851 ATAGCTTAGC CCCATTTCTG CCCCACTGTG TGTAGCTCAG AGTCACCTTG
140901 TTACTCTAGA GCCAAATTCA TCACTGTTTA GGTACCCACA TTAGAAAAGA
140951 GTCAAGTGTT GGCAAGGGAA TTCCAATCAA GCCACAAGCC TGGAAAAGGA
141001 GCTCTCTATT CTGAGCTCTC TGAGTTCTCT ATTCTGTTTA ATTGGTCTAT
141051 GCGTCTGTCG TTGTACCAGT ACCATGCTGT TTTGGTTACT GTAGCTTTGT
141101 AGTATAGTTT GAAGTCAGGT AGTGTAGTAG TGTAATAATG CCTCCAGTCT
141151 TTTTTTTTTT TTTTTTTTTT TTTTTTTTTT TTTTTTTTTT GCTTAGGATT
141201 GTCTTGACTA TTCAAGCCCT TATTTGGTTC CATATACATT TGAAAATAGT
141251 TTTTTTTTCT AATTCTGTGA AGAATGCCAA CAGTCATTTA ATGGGAATAG
141301 CATTGAATCT ATAAATTACT TTAGGCAGTA TGGCCATTTT TATGATATTG
141351 ATTCTATCTG GGAACCTGGA ATGTTTTTCC ATTTGTTTGT GTCCTCTCTG
141401 ATTTCCTTGA GCAGTGGTTT GTATTTCTCC TTGAAGAGGT CCTTCATTTC
141451 CCTTGTTAGC TATATTCCTA GGTGTTTTAT TGTTTTGTAG CAGTTGTGAA
141501 TGGGAGTTCA TTCATGATTT GTCTCTCTGC TTGCCTGTTG TTGGTGTATA
141551 GGAATGCTAG CAATCTTTGC ACATTCATTT TATATCCTGG GTTTCAGTAT
141601 TTTAAAAACT TACTTCAGGT GATTCTATGT GTGCAACCAT GATTGAGATA
141651 CACTGTTATA GAATCTAGGA TGTGATAAAC TAGAAGAACA TAACTAAAGT
141701 TTTGCATTTT TCGGGTGTCT CAGTTTCCTC ATTTATAGAT GGAGTTGGTA
141751 TGTGTACCAA GTTCATAGGC TTGTTCTGAG TAAATTAGTG CATGTAAAGT
141801 GCTCCACAGA ATGTTAGCTG TTGTGATGCT TTACTTTCCA TTGCACTTCC
141851 TGACTCCTAG CCTTTCTTTT CCTTGGCTCT TTTTATGCTC ATGTCAGATG
141901 CCTCTATTGT TTCTTTCCCC CCAGAATATC CTCCACTTTA TCTTGCTCTG
141951 CTCAACATCT TTAAAGTATA GAATCAACAG ACTGCCATGC CACCCAGTCT
142001 GTCTGACAAT TGAGGCAAAT TCCCTAAGTC CTCTTGTTCT CCTTCTGAGA
142051 TTTCCACCTG CTCTAACCCC TTCCAATATT TCAGATGCCG TCTCCAGCTA
142101 TGATAATTTA ATCAGTGTTT GCTCTGCTCA TCCTTGATAT GTGAGTCCTA
142151 AGATTTTAAG CGATCATTTC CCTTCTAAGT CATGTATGAC CCATTAGTCC
142201 CTCCATTCTT TTTTCTTACC CCTCATTTCA TATTCTCTTT ATGGCTACTC
142251 CTGTTGATGT ATCCATTTGG CCACACTTCT TAAACTTCTC CACCTAAAGC
142301 AGAGGAAAAA GAACAAGTTG AACATGAACC CTTTAAGGGG AATGGGGTCT
142351 GAAGTGTCAC ACTAAAAGGT CATCTGCAAG TATGTATTTC ATATCTTTGT
142401 TTAAATAAAA TAGTTACATA GTAGAGGGAA AAAAAATCCA TGTGGATTTT
142451 GCATTTCACT CAATTATAAC CTTGATTTTT AATGCTAAAA ATTATTTTTC
142501 CTAAAATCTT GGGGTAAAAG TGTTGCTCCA AAGAGCTTTT ATCAGATTAT
142551 GTTTATCCTG TAGCTGCCTG TCCCCTGTGA CCGATACTGG AAACCCTCAG
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FIGURE 3TT

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142601 GATTACAAAT GCCTCCGTTT GCAAGTAAGA GTGAAATACA GCAGAACTGT
142651 GTCTTCTCCT TTGTCTTGTT CCCCATCTCT CTTCTGTGCT TTGTATTGTT
142701 TCCTCTCCTG TCACCTAAAC AGGCACTCTG AAAGAAAACT CTCCAGTACT
142751 GGAGAACTTA GCATATTCTA ATTCCTAGGT TAAAAAAAAA TAATAAATGA
142801 CTGAATGATT TTTTTTAAAG AATATTTTCC ATCAGAAGAA ATTTGGAAGT
142851 ATTTTGTTGC AGAATTTTAA AACATTTGAT CTGGGTCTAA TTCTGTCCTG
142901 GGACTGGTAA TCATCTTTTT TTGAGGCTAA ATTTTCTCAT TTTGATGAAA
142951 AAGTCATCAA TAGATGTTGA AAGCTGGACA GTGCAGTGTC AAAGCAAATG
143001 CTTTGCATGT CTGCAAGAAA GTCACAAATA AAGAAGGCTC TGCTGACTAA
143051 AAGAGAAAGA TACTTAATCA ACTCCAGTAC CATTGTTGAG GGGAACATTC
143101 TATCAGGATT CAGTATAGAG AGATATTTTT AGGCTATTCA CAAAATCCAG
143151 GTAGAACCTC CAAGCTACAT TTACAATAAT ACTAGCTTTT AGATTAATTG
143201 TTGTTTTTTA AATATGTATT AGCCTCTTAT ACAAATATAA GGAGTTACAA
143251 ATTATTATTA CAATAATCTT GGCTTTCGTG ATTGTCCAAT GTATTTACAC
143301 GTACCGAGAG CTTTATTTCT CCGTATAGTT TCAAGTTACT GTCTCGTGTC
143351 CTTTCATTTC ACCTTGCAGG ACTCCTTTGA GCATTTCTTA CAGGGAAGTT
143401 CTAGTGGTAA TAAACTCCCT CCACTTTTAT CTGGAAACAT CTTAGTTTCT
143451 CTCTCACTTT TCAAGAACAG TTCTGCCAGA TAGAGGACCC TTGGTTGATA
143501 GGTTTTTTTC TTTTAGCACT TTGAATATAT CAGCCCACTG CCTTCTGGCC
143551 TCCAAAGTTT CTGATAAGAA ATCTGCCCGT CATCTTATGA TGTACTTGAC
143601 AAATTTTTTC TCTCTTGCTG CTTTCAAGAT TCTCTCCTTG TCTTTGGCTT
143651 TAGAAAGTTT GCTTATATTG GCTGGACATG GTGGCTCACA CCTGTAATCC
143701 CAGCACTTTG GGAGGCTGAG GCAGGCGGAT CACTTGAGGC CAGGAGTTTG
143751 AGATCAGCCT GGCCAACATG ATGAAACCCC TGCCTCTACT TAAAATTCAA
143801 AAATTAGCTA AGTGTAGTGG TGCACACCTG TAATCCCAGC TACTTGGGTG
143851 GCTAAGGCAA GAGAATCTCT TGAACCCAAG AGGAGGAGGT TGCAGTGAGC
143901 TGAGAGCATG CCACTTCACT CCAGTCTGGG CAACAGCA AAAGTCTGTC
143951 AGAAAAAAAA AAAAAGGAAA GTTTGATTAT ATTATGTGTC AATGTGGGTC
144001 TTTTTGAATT CATCTTACTT GGGATACACT GTGCCTTTTT GGATTTGGGG
144051 GCTCATGCCT TTCAGCTATG ATTTCTTTAA GTATTCTGTT TTCCTTTTTC
144101 TCTCTCTTCT CCTCCTGGGA CTTCCACAGT ACGTACACTG GTTTGCTTGA
144151 TGGTGTTCCA TACATTCCTG TAGGCCAGGG ATGTCCAATC TTTTGGCTTC
144201 CCTGGGCCAC GTTGGAAGAA GAGGAATTGT CTTAGGCCAC ACATAAAATA
144251 CACTAACACT AACGATAGCT GATGAGCTAA AGAAAAATCA CCCTCAAAAA
144301 AATCTCCTAA TGTTTTAAGA AAGTTTACAA ATTTGTGTTG GGCCACATTC
144351 AAAGCCATCC TGAGGCACAT GTGGCCCATG GGCTGTGGGT TGGACAAGCT
144401 TGCTATAGGC TCTGTTCATT ATTCTTCAAT CTTTTTTCTT TCTGTTCCTC
144451 AGACTCAGTA ATTTCCACTG TCCTGTCATC AAGTTTGATA CTGATTCCTT
144501 CCTTGCCTGC TCAATTTTGC CGTTGAAACC CTGTAGCAAA TTTTTAAATT
144551 TTAGTTATTG CACTTTTCAG CTCAAGAATT CCTTTTTAGT TTCTTTTTAG
144601 GTTTTCTATA TTTTTATTAA TACTTTAGTT TTGTTTGCAC ATCATTTTCT
144651 TGATTTTCTC TATATCTTCC TTTAGCTCTT TGAGCATCTT TAAGATAGTT
144701 GTTTTGATGT CTTTATCTAG TAGATCTACT GTTAGGTCTT TTTAAGGGAT
144751 AGGTTTTTTG GTTTATGTTT TTTACTGTGA ATGAGCCATA CTTCTCTATT
144801 TCCTGGCATG CCTTGTTATT TTTTGTATTG GACACTTGAA TCTAATAATG
144851 TGATAAATCT AGGAAAATCA GATTTCTCCC ATCCCCAGGG TTTGCTGTTT
144901 TTTGTTATTG TTTTTATTTT TATTTTTTAT TATTGTTGTA AGCTGTCTCC
144951 ATGCCAAGGA TCAGCTGAGG TGTAAACATA AGATCTTCTT AGGTCTTTTC
145001 TGAGCCTGCA CCCTTCCCTG GTCATGTGCA GTCACTTTCT AATTTTCCCT
145051 ACACATGCAG TTGTTTTTGA ATGTCCCAGC CTTTCACGTG TGGCTCCCAA
145101 AAGGAGGAAA GGAGAAAAAT GAAGAGGGTG AAAAGGTGCT GGCCCTTTAA
145151 TTCTCCCAGA AGTCACTTCA GCCTGAGGGA GAGTGGCTGG CAACATTGTG
145201 GGGGAGGTGC AACAACAATG GCCATCAAGC ATTTTGTTTG CACCTCTGTG
145251 ATCAGAAGCA GCAGTGTCGG AAGCACAGAT CCTCAGAATT TGGAGAACAC
145301 AGTTCTTGCT TTCCACCCTG ACTCTCACAG GCTGTGTGCA AACTGCTCCG
145351 GAACATGTGT GTGCTCAGCT CCCTCCCATG GGGCTGGAGG ATGAGGGATG
145401 GGTAGCTGCT GCTGTGCTAA GAGCTTAAGT TGGTCATAAT TAACTGCGCT
145451 TTGCCACCCA AGCCTTCCCT GAAAGTTGCA AGCTTTCAAT AGACTCCAGA
145501 GTTCTAAAAT AGTGACATTA GACAGATTCT GCCAGTGCAA TCGCTGTCTA
145551 GGAGGGGAGA CAGATTCCTG GTGCTTCCTG TTTTGCCAGC TTCCCGGAAT
145601 CTTCTTCACA TAGCATCCAT TTTGAAGATA CTACTTACTT CTCAATTTGG
145651 GGCTATTCAT TGAATAGACT GTCACCAGGT TATTGGCTGT TTGAAGATTC
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FIGURE 3UU

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145701 TCATTTGTCT GCTAACTATA CCTCTATTTT TTTTCTACGT TCACCTGGAA
145751 GACATGTCTT CTTCAAGAGC ACCTTGACTC TGTCCAGAAG GAGTTCATAA
145801 TTTTCAACAG AGAAAAGTAA GTAATTCCTG GGAGAACAAC AGCCCCAGAA
145851 ATGGTGGCAT GTTTCAGCCA GACTTTACTT GCAGAGAAAA TATATTTTTA
145901 ACATTTTAAA AATTATTTTC TAATTGGGAA AATGATGCAA TCTATTATAG
145951 AAAATGTAGA AACCTTTTTT GTAAGGTATT TAACATTTTT TAATTGATAA
146001 ATTAGCCTAG CATCAAGTTT TTGTTTGTGA GAAGGGAAGA GGAATTAGGA
146051 TTTAAACACT TAAAAATCAA AGCCTTTTAA AAGATTTCCT TGGCTCATGC
146101 TTATTTATAA ATTATTGGGC TTAATATTAT TTCAAAAGCT TAAACCTTTC
146151 ATTTTATTTT TCAAAGAATA AAACATCTTT TTTTTTCTTT TCTTTTTAAG
146201 AGTAAACAGG GACTTTAACA AAAGACAACC AAATCTAGCC TTGGAACAAA
146251 CCAAAGACCC ACAAGGTGAG GATGGTCAGA ATAACAACTT GTAAAGGCCT
146301 CATGTCTTCT TCTTGGGACA ATCTCATGCC AGAAACTTCT AATTACATAT
146351 GTCAAGAAAA GCTGACAGTA GTTCTTGCCA CTCCACACAC CATGACTTAG
146401 AAAATGTGAA TGAATATATT TCAAAAAAGG CAGCACAACA CAGTGAAGGG
146451 TCCTGGGCCT GAGCTCCTGG GATGTCATTT CACATCAATC AACTGTGTGA
146501 TCTAGAGCAA GTCACTTAGC CACTTTCTGT GCTTTACTTT ATTTATCTAA
146551 AATGAGAGGG TTATACTAGA CGAGCCATAC CCTGCCTTTT TAGTGCTATA
146601 GTTGTTATTC TAAACCGCCT TTATTTTTAT TTTAAAATTA ATATATGAAT
146651 ATAGATTTAT TTTTCCACTC CTTCTAATTA TGCAGTGACA AATGGACAAA
146701 TGGACACAGG ACTCAGTGAG ACTTTTCAGA CCTCGAAAGT TTCATAAAGT
146751 GGTCAGAATG CCCCAGGCTA CTTGGATAAA GATAAGGAAT TCTATCAGGG
146801 AGGCATGAAT GGAATCAGAT TAAAAGTAAC AGAGATGGAT GAGGGCCTTC
146851 CAGTGATATG CGTGAATCAG CATTAGATCC GCTTATCTCA GCTGGCAGGA
146901 GCCTGCTGTG CACACCACTT CCCAGCTCCC TCTTCAACAA TGTGAAAGTG
146951 GTAACTTGAA ATTGGTAATA ATGGGAGCAT TTACACCACG GAAACTGGTA
147001 AATGCTCGTT TTTTCCCTCC TAACAAGTGA ATTGCTAAAT ATTAGCCCAC
147051 CACTCCTTCC AAGAAGCATG TTCCTTGAGG GCTAATTGTC CTCTGAAGAT
147101 TAGCAGAGAC CTGTATCTGG AGAGGATCAG AAAAGAATGT CATCACACTG
147151 AAAGTATGTC CACCTTGCAG TTCAGAAAAG TTGCATCTTA TATGGGGTTT
147201 ATTGTCTAAG TTAGAAATGA ATTTAGAAGA TAGTAAAATT TACCGTTGAA
147251 AAACCCCTTA AATTACCCAT AAAGTATATG GGAAGTATCT TTTCTCAGTA
147301 AAGCCCAATA CAGTGTCACC TTTCACTAAT GAAACAAGCC ATTGCTTTTG
147351 TTTTGTTTTG ACTTAGTTAT TTTTATTTTT GGTCTCATTT TGGCTAATAC
147401 CAGATGAGCT AAAATGTTGA ACAAATTATA CTTGTTTTTA TAGACTAGAA
147451 TTACTCTTTT TTTTCTTTTC AGGCAGAGTC TCACTCTGTC ACCCAGGCTG
147501 GAGTGCAGTG GCATGATCTC TGCTCACTAC ATCTGCCTCC CGGGTTCAAG
147551 TGATTCTTGT GTCTCAGCCT CCTAAGTAGC TGGGATCGCA TGTGTGTGCC
147601 ACCATGTGTA GCTAATTTTT TGTATTTTTA GTAGAGATAG GATTTTGCTA
147651 AGCTGGCCAG GTTGGTTTCA AACTCCTGGT CTCAAGTGAT CCGCCCACCT
147701 TGGCCTCCCA AAGTGCTGGG ATTACAGGCG TGAGCCACCA AACCTGGCCT
147751 TAGAATTACT CTTAGAACAG TGGAATGCCC ACACATCCAA GACAGGCAAG
147801 TTCATGGAGA CTAAGGGAAC AGTGGTATCA TGTCTCCCTT CTCCCTTGTG
147851 CTTACTACAA GAATGGCAGG CAGAATTCCC TACTTATTTA AAATATCACT
147901 GATGTCTCAC TCTTTTTCTT TATATTTTAT TTATTGATTT GCCACAAAGT
147951 TTAATTCACC TAAGTGAGAC GTGCATATGA TGTAACTCCA CTGTACAGAT
148001 ACACAGATCT TTACAGAAGA ACTATTTTTG GCAACCCCTA TGCCCCTGGG
148051 TAGGGTCCAG AAGTGAACAG GCTTGGTGGG GGATTGTTTT CACCTCTTGG
148101 CTACTCAGAG TACCTAAACC TGTCCTTACT TATGGAGAGC ATGTGTCACA
148151 CCAAGATGGC AGTAAGCTGG CAACTGCGAA GACCTGACTG ATGCCCATTT
148201 GGGAAGCCAG GCAAGTGAAA ATGGACCGAA GAAACAGAGA TGGCTGTCTT
148251 TTATGCAGGG CTTTTCCATA AAGAGGTTAC ACTGGGGCAA CCAAGTATGT
148301 GTAGAAAGCC AGAGCTAAAC TTCAGCTTGG CATTCACAGT TTTCTCTTCA
148351 CTGAGCTAAT AGGCCCAGAG TTTCGGGCAG AGCTGTGAAA TAGTGCTTCT
148401 CTAATAGCAA CCATATTATT GTTACATAAT TAAAAGCCAG CTCTTTTGTT
148451 GTTTGTTTGA TTCCTTTTCC CTACAGTTCC CACATCATTT GTCTGTGCTA
148501 TTCTGTTTTT CTCCAAACAC TATAAACTTG AAGCAATTGC CCTGACTCGA
148551 TTTCAGAGAA GGGGATG
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(SEQ ID NO: 3)
FEATURES:
Start: 2003
Exon: 2003-2054

FIGURE 3VV

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Intron:   2055-22564
Exon:     22565-22567
Intron:   22568-39943
Exon:     39944-39999
Intron:   40000-41067
Exon:     41068-41219
Intron:   41220-79497
Exon:     79498-79500
Intron:   79501-86044
Exon:     86045-86218
Intron:   86219-105152
Exon:     105153-105190
Intron:   105191-113211
Exon:     113212-113301
Intron:   113302-123667
Exon:     123668-123765
Intron:   123766-132805
Exon:     132806-132922
Intron:   132923-133775
Exon:     133776-133848
Intron:   133849-134127
Exon:     134128-134155
Intron:   134156-134530
Exon:     134531-134545
Intron:   134546-135588
Exon:     135589-135601
Intron:   135602-137397
Exon:     137398-137526
Intron:   137527-145751
Exon:     145752-145816
Intron:   145817-146200
Exon:     146201-146291
Stop:     146292
```

SNPs:

DNA Position	Major	Minor	Domain	Protein Position	Major	Minor
210	G	A	Beyond ORF(5')			
332	T	A	Beyond ORF(5')			
1131	T	A	Beyond ORF(5')			
1221	C	T	Beyond ORF(5')			
2011	G	C	Exon	3	A	A
4309	A	C	Intron			
4345	C	T	Intron			
4651	T	C	Intron			
5037	A	G	Intron			
5126	C	T	Intron			
6048	G	T	Intron			
6229	–	T	Intron			
6328	G	A	Intron			
6350	C	T	Intron			
6382	G	T	Intron			
6434	A	–	Intron			
6722	A	G	Intron			
6751	C	T	Intron			
6752	A	T	Intron			
7070	T	C	Intron			
7306	T	C	Intron			

FIGURE 3WW

7339	A	G	Intron
7531	G	A	Intron
8902	A	G	Intron
9471	G	A	Intron
10023	C	T	Intron
10594	C	G	Intron
11233	C	T	Intron
11295	–	A T	Intron
11534	–	T	Intron
11757	T	C	Intron
11951	G	A	Intron
12901	C	A	Intron
13040	C	T	Intron
13081	A	G	Intron
13173	G	T	Intron
13272	C	T	Intron
13333	A	– G	Intron
13485	C	A	Intron
13933	A	T	Intron
14086	G	A	Intron
14094	C	T	Intron
14141	G	–	Intron
14831	T	C	Intron
15319	T	C	Intron
15321	T	C	Intron
15335	A	G	Intron
15477	G	A	Intron
15650	T	C	Intron
15880	C	T	Intron
16944	G	A	Intron
17061	C	T	Intron
17494	G	C	Intron
17642	T	A	Intron
17737	A	C	Intron
18068	A	G	Intron
18339	C	T	Intron
18361	C	T	Intron
19218	A	G T	Intron
19298	C	A	Intron
19629	C	T	Intron
19679	G	A	Intron
19981	A	G	Intron
20014	C	T	Intron
20280	C	T	Intron
20612	A	C	Intron
21966	C	T	Intron
22017	T	C	Intron
28009	G	A	Intron
28059	T	A	Intron
28580	T	C	Intron
28595	A	C	Intron
28823	–	A C	Intron
28827	C	G	Intron
28842	G	T	Intron
30128	T	A	Intron
30150	T	G	Intron
30188	C	T	Intron
30453	T	C	Intron
34990	A	G	Intron
35203	G	A	Intron
36206	G	A	Intron
39692	C	T	Intron

FIGURE 3XX

40095	A	G	Intron
40191	T	C	Intron
40287	G	A	Intron
40384	T	C	Intron
40510	G	A	Intron
41664	C	T	Intron
48324	T	G	Intron
48423	C	T	Intron
50015	A	C	Intron
50095	T	G	Intron
52300	A	G	Intron
52623	C	G	Intron
52773	G	A	Intron
53140	G	A	Intron
53848	A	T	Intron
57636	–	A	Intron
57693	A	T	Intron
58585	T	C	Intron
58649	T	C	Intron
62188	A	T	Intron
63478	G	A	Intron
65457	C	A	Intron
69947	A	G	Intron
69981	C	T	Intron
71165	G	A	Intron
71347	A	G	Intron
71903	A	G	Intron
71908	C	T	Intron
71994	G	A	Intron
72010	T	C	Intron
72612	G	A	Intron
73294	A	G	Intron
73385	C	G	Intron
74121	G	A	Intron
75646	A	G	Intron
75698	C	T	Intron
79007	–	A	Intron
80043	A	G	Intron
80499	G	C	Intron
80940	A	– T	Intron
81615	G	T	Intron
82599	C	–	Intron
82952	–	G	Intron
85020	A	T	Intron
88843	C	T	Intron
89700	G	A	Intron
90002	G	A	Intron
90615	A	G	Intron
92506	A	G	Intron
92558	T	C	Intron
92667	G	A	Intron
92803	A	T	Intron
95079	T	A	Intron
95089	G	A	Intron
96495	G	A	Intron
97070	T	A	Intron
99913	A	C	Intron
102375	C	T	Intron
102686	–	A	Intron
102687	A	C	Intron
102939	C	T	Intron
106162	G	T	Intron

FIGURE 3YY

106378	T	G		Intron
107310	C	T		Intron
108663	C	A		Intron
108876	A	T		Intron
110733	C	G		Intron
111546	A	G		Intron
116728	T	C		Intron
118403	G	T		Intron
118491	C	G		Intron
118888	A	G		Intron
125444	–	A	T	Intron
125810	T	C		Intron
126092	T	C		Intron
127506	G	A		Intron
127878	G	T		Intron
139738	T	C		Intron
140261	C	T		Intron
141590	T	G		Intron
142613	C	T		Intron
142774	C	A		Intron
143288	G	A		Intron
145610	A	C		Intron
148360	T	C		Beyond ORF(3')

Context:

DNA
Position

210
TCCCTCTCTCATACCATTTAATTGGTTGCTTCCTAATTAATGACTCTCTTTGCTCTCTAT
TTAATGATTCTTGCTAAAGTCCATAAGGCACTTTGCCAGCAGTTGGTTTTTAGTATGAAA
AGTAGCATTTCCTTAATGAGTCTGAGTCTGCCTTCCAAATGAAGGGTTTACTTACATTTT
CCTAATGGGAAAACGAGCTTTTCTTCTAC
[G,A]
CTTCCTTAGGGGTTTCATAAGTTCTTTTTTCAATAACTCATCCTTAACACTTTCTCCAATT
CTGCCTGTAATCAATATTCCCTTCACATGTAAAGAGCTCAGGAGGAAATCAACTATTTTT
TTAAAAATACGCAATAAGGAAATTCTGCTACTCTTAGAAATAGCAGGAGCTAACATTCAT
TCTTTGCATATCATGTGCTAGGCATTGTGCCAATTACCTTATATACATTGTCTCATTATA
TGTATCCATGACCATATATGTGCTAAGCATGAAATTTTCTTAAGCCAGATAGCTGAGTAG

332
CCTAATTAATGACTCTCTTTGCTCTCTATTTAATGATTCTTGCTAAAGTCCATAAGGCAC
TTTGCCAGCAGTTGGTTTTTAGTATGAAAAGTAGCATTTCCTTAATGAGTCTGAGTCTGC
CTTCCAAATGAAGGGTTTACTTACATTTTCCTAATGGGAAAACGAGCTTTTCTTCTACGC
TTCCTTAGGGGTTTCATAAGTTCTTTTTTCAATAACTCATCCTTAACACTTTCTCCAATTC
TGCCTGTAATCAATATTCCCTTCACATGTAAAGAGCTCAGGAGGAAATCAACTATTTTTT
[T,A]
AAAAATACGCAATAAGGAAATTCTGCTACTCTTAGAAATAGCAGGAGCTAACATTCATTC
TTTGCATATCATGTGCTAGGCATTGTGCCAATTACCTTATATACATTGTCTCATTATATG
TATCCATGACCATATATGTGCTAAGCATGAAATTTTCTTAAGCCAGATAGCTGAGTAGAA
TTTTAAAATATTATTTTGTACAAAATCTAGACCTTTACCCCATTTGGGGGATAGATCTGA
AGATCTGGGCTCATGTTTCCATGTGGTGACAATCTGTTTGATCTGAGCACAATTACTTTA

1131
AGGACTGGAACACAGGATGCTGCCTCTCTTTACCATTATGTTTTAAAGTGGAGCAAAGCC
GTAGTTTTCAGGATCTTTTCTTGTTCACACATATCATTTAATTTGAGCCTCAGAGCGGCT
AACAGTTTTGAGCACTTATGCTATGAAAATGTTTTGTGTATTCAGTTAAATGTATGCATA
TCATACATTTATGTAACTCAATACATATATATAAATGTGATATAACATACGTATGATATA
ACAGAGTTATATATATGTGTATTATTTAACTTAATATATAATGAGTTAAGTGTATGCATA
[T,A]
CATAGATTTATGTAACTCAATATATATAAAGAGTTATATAATACAACAGAGTTGATATATAT
ATAAATGTTGTATATAAACATAATATATACGTTAATATATATTAACAAAGAGTTGTATAA
TACAACACAGAGTTAATAATATATAAATACAACACAAAGAGTTATATATGTGTGTATTAT
ACATTTAACTTAATATATAATGAGTTAAATGTATGTCTGTCCCATTCAACTCTCCATTGA

FIGURE 3ZZ

GGAAAGTACCATTATCTTCCCCAAGTTCAGAAGAAGAAAACAGAGAAATATATTGAAATT

1221   ATATCATTTAATTTGAGCCTCAGAGCGGCTAACAGTTTTGAGCACTTATGCTATGAAAAT
GTTTTGTGTATTCAGTTAAATGTATGCATATCATACATTTATGTAACTCAATACATATAT
ATAAATGTGATATAACATACGTATGATATAACAGAGTTATATATATGTGTATTATTTAAC
TTAATATATAATGAGTTAAGTGTATGCATATCATAGATTTATGTAACTCAATATATAAAG
AGTTATATAATACAACAGAGTTGATATATATATAAATGTTGTATATAAACATAATATATA
[C,T]
GTTAATATATATTAACAAAGAGTTGTATAATACAACACAGAGTTAATAATATATAAATAC
AACACAAAGAGTTATATATGTGTGTATTATACATTTAACTTAATATATAATGAGTTAAAT
GTATGTCTGTCCCATTCAACTCTCCATTGAGGAAAGTACCATTATCTTCCCCAAGTTCAG
AAGAAGAAAACAGAGAAATATATTGAAATTCAGCAATTTGCTGGTGTGGTCAAGTCCAAC
CCAGAACTTGCTTCTTTTACATTGTAGTACCCTCCAGGGTATGCAGAAACAGATAGCTAG

2011   TAGGTCTGCTATTGATGAGCCATGCAGTGTTTTCTCCTGTTGCTTGATGTTTTTATTCTG
AAATCATGGTTGGTTTTCAAACACAAAGTTTTCACTACAGTGATACAGATGAGGTTTAT
GTTTCCGCCACAGTCTATACTCAGGGTGCCTAGAGTATAGCATATTATTAGGGTACTATT
TCTTTTCCTATCCTAGATATCCAACTAAGGCTTCGGGACATGTTTTGAGCGAAGATGGGT
GTTTCTGCCCGGATAGTATAAATCGAGGATCCAGGTCTGGGCAGATTCAACCATGGGAGC
[G,C]
AACACTTCAAGAAAACCACCAGTGTTTGATGAAAATGAAGATGGTAAGAAATATGGGATA
GTGGCATATAAAAAATAGAATTTTGCAAAATTCAAGTATATGCTTCTAGTTTTCATAAGTT
AAGCATAAGCATGGTCTGTAGGGCCTTGAAGGAAAAAGGCAAAGCTGCATGAGTGAGTCT
GAGGACTTTGTAGGCTCATAGCTAGGTTTTACCTTCCACTTTCCATGGGACCTTTGGCAG
CTTTCCTAATCTCCACTATACCAATGTCCTTTGTCCAAAGGGAGCTGCAGTTGGGCATGT

4309   GAAAACTTTTTTTTTCATGGCTATTGTGATTGCCTTGCTTTAACTTATCAAATAGTAAAAG
CAAAGATCTAGAGACTAGTGATATTACTTAATTTTTCTGTCTCTAAAATGGAAAGACAAA
TAGGCTTGCTTTTCATTTAGTTGGTTTCCTCTGCTTCCTCTGGACTCAGAGCTAATGTTG
TACATGAGGCTGGTCGTCAGAGAATAGGGTGGAAAAGAGAGGCCAGCTGCATACTTTTAA
CTTGCTGGGCTACATTTGAAGGTAGTAGAATAGCATTATGATGAGAAAACACAGAAATGC
[A,C]
TAACTCTTCCTTGATTCAGCCAGGCTTTGTTCTTGCGGGATGCCCAAGAAAGCTACATAA
CCAAAGAATTGTGACAATTGGGAAATAAGATACCCCTTTTTAGTTACTTTAAAGGACTCT
AGAAAAACTAGGTTGAAGGAGAGTTAGGCTTAGGGACCAGACAGGTCTTTCTTAACACCC
TCTAGGTCACCACCTTTTCTGTTGTCTGGCTTCTCAGCCCAATGAGATGAACCCACTGCA
GCACCCATAAAGGAAAGATCTGAGCATAGCAACAAGTCTGTGCCTCCCAAAGGTGCTAGG

4345   CTTTAACTTATCAAATAGTAAAAGCAAAGATCTAGAGACTAGTGATATTACTTAATTTTT
CTGTCTCTAAAATGGAAAGACAAATAGGCTTGCTTTTCATTTAGTTGGTTTCCTCTGCTT
CCTCTGGACTCAGAGCTAATGTTGTACATGAGGCTGGTCGTCAGAGAATAGGGTGGAAAA
GAGAGGCCAGCTGCATACTTTTAACTTGCTGGGCTACATTTGAAGGTAGTAGAATAGCAT
TATGATGAGAAAACACAGAAATGCATAACTCTTCCTTGATTCAGCCAGGCTTTGTTCTTG
[C,T]
GGGATGCCCAAGAAAGCTACATAACCAAAGAATTGTGACAATTGGGAAATAAGATACCCC
TTTTTAGTTACTTTAAAGGACTCTAGAAAAACTAGGTTGAAGGAGAGTTAGGCTTAGGGA
CCAGACAGGTCTTTCTTAACACCCTCTAGGTCACCACCTTTTCTGTTGTCTGGCTTCTCA
GCCCAATGAGATGAACCCACTGCAGCACCCATAAAGGAAAGATCTGAGCATAGCAACAAG
TCTGTGCCTCCCAAAGGTGCTAGGCTCTCTGTCTGTTTATGCAGACAGTTGCAAGGCAAA

4651   GCCCAAGAAAGCTACATAACCAAAGAATTGTGACAATTGGGAAATAAGATACCCCTTTTT
AGTTACTTTAAAGGACTCTAGAAAAACTAGGTTGAAGGAGAGTTAGGCTTAGGGACCAGA
CAGGTCTTTCTTAACACCCTCTAGGTCACCACCTTTTCTGTTGTCTGGCTTCTCAGCCCA
ATGAGATGAACCCACTGCAGCACCCATAAAGGAAAGATCTGAGCATAGCAACAAGTCTGT
GCCTCCCAAAGGTGCTAGGCTCTCTGTCTGTTTATGCAGACAGTTGCAAGGCAAAGGAAG
[T,C]
AGGAGGGCAAGTCCACCTACTATAAACCTGTCACTCTCTAGACATGAAGAATAGAGGAGG
AAACAAGTTGGTCCTTGCTCTGTCATTGTGAACCCCATGTTCTGATGATGGAAGGCTGAC
AATAAAAAGGTAAATAATACATAAACCAGATAATTTCACAGTGCCTTAAAGTGCCACCAA
GGAAATGACTCCTAGTGATCTTACAGACAGTGACAGTGATGGTGAGGAGGCCACTTTAGA
TAGGGTGGCTGCGGTTGTCTTTCTAAGGAGGTGACATTTGGGCTGAAGCCTGAAAGATGA

FIGURE 3AAA

5037   TTGTGAACCCCATGTTCTGATGATGGAAGGCTGACAATAAAAAGGTAAATAATACATAAA
CCAGATAATTTCACAGTGCCTTAAAGTGCCACCAAGGAAATGACTCCTAGTGATCTTACA
GACAGTGACAGTGATGGTGAGGAGGCCACTTTAGATAGGGTGGCTGCGGTTGTCTTTCTA
AGGAGGTGACATTTGGGCTGAAGCCTGAAAGATGAGAAGAAGCCATCTATGAAATGACAT
GAAAAGAATAGTTCAAGAACAGGAAAAACAAGTCCAAAATCCAAATAATGACAAAATCAG
[A,G]
ATTGAATAGTTGCCTATATCTTAACGTTCTCTCATGAGCACTAGTTTGCCAAAGAGACTG
CATTTATTGCCATGTTAACTTATTTCTTCAAAAGATGATTGATTTGAGGAGAAAAAGTAT
GCCATTCTAGGGAATTTACTTTGCTTTAAAAATTCAGTACATTTTGTAAAGTTCATTTGAC
TCTTCACATAAATCTGGATTGAGCACAAGGTAAAATTGTATCTGATTGCTGTGAAGCTCC
TGACCAAGAAAAAGCAACCAAAAAGCACTGATTAACCAAACAACATTAATGCTTATGTCA

5126   CACCAAGGAAATGACTCCTAGTGATCTTACAGACAGTGACAGTGATGGTGAGGAGGCCAC
TTTAGATAGGGTGGCTGCGGTTGTCTTTCTAAGGAGGTGACATTTGGGCTGAAGCCTGAA
AGATGAGAAGAAGCCATCTATGAAATGACATGAAAAGAATAGTTCAAGAACAGGAAAAAC
AAGTCCAAAATCCAAATAATGACAAAATCAGGATTGAATAGTTGCCTATATCTTAACGTT
CTCTCATGAGCACTAGTTTGCCAAAGAGACTGCATTTATTGCCATGTTAACTTATTTCTT
[C,T]
AAAAGATGATTGATTTGAGGAGAAAAAGTATGCCATTCTAGGGAATTTACTTTGCTTTAA
AATTCAGTACATTTTGTAAAGTTCATTTGACTCTTCACATAAATCTGGATTGAGCACAAG
GTAAAATTGTATCTGATTGCTGTGAAGCTCCTGACCAAGAAAAAGCAACCAAAAAGCACT
GATTAACCAAACAACATTAATGCTTATGTCATTTTTGATATCCATATTTTTATATACATA
ATCATAATGTATAATCAAACTGGGCCAGTATCAAGGGCACTAAAATGAGCCAACTTAATT

6048   ATCTGATCTTGAACACATAATTTTATTAGTTACTTATGTTGATCTTTATTCAGCAAAAAC
AAAGTAGGAGATTTTCAGGCTAGGCATGGTTGCTTACGCCTGTAATCCCAGCACTTCAGG
AGGCCGAGGCGGGCAGATCACGAGGTCAAGAGATCGAAACCATCCTGGCCAACATGGTGA
AACCCCATCTCTACTAAAAAATACAAAAAAAAATTAGCTGGGCATGCCAGTGTGCGCCTGT
AGTCCCAGCTATTCAGGAGGCTGAGGCAGGAGAATCTCTTGAACCTGGGAGGTGAAGTTT
[G,T]
CAGTGAGCTGAGATTGCTCCACTGCACTCCAGCCTGGCAACAGAGCAAGACTCTGTCCAA
AAAAAAACGGCTTGCTTATTTGATTATATAAGATATCTTTCATAAATTAGATCTCAAATT
ATACTATTGTTTTGCAGTTTTAGCTTTTATGTTTTAGGGCAAATCTTAAGTCCTAATTAC
TTTTTTTTTTATTATTGTGGTAAAATGTATATAACAAAATGTACCATTTAATCATTTTAGA
ATATACGGTTTATGACATTAAGCACATTCACGTTATCATGCAACCATCACCACTACCCAT

6229   ACCCCATCTCTACTAAAAAATACAAAAAAAAATTAGCTGGGCATGCCAGTGTGCGCCTGTA
GTCCCAGCTATTCAGGAGGCTGAGGCAGGAGAATCTCTTGAACCTGGGAGGTGAAGTTTG
CAGTGAGCTGAGATTGCTCCACTGCACTCCAGCCTGGCAACAGAGCAAGACTCTGTCCAA
AAAAAAACGGCTTGCTTATTTGATTATATAAGATATCTTTCATAAATTAGATCTCAAATT
ATACTATTGTTTTGCAGTTTTAGCTTTTATGTTTTAGGGCAAATCTTAAGTCCTAATTAC
[-,T]
TTTTTTTTTATTATTGTGGTAAAATGTATATAACAAAATGTACCATTTAATCATTTTAGAA
TATACGGTTTATGACATTAAGCACATTCACGTTATCATGCAACCATCACCACTACCCATC
CTCAGAACATTTCTCTTCTCGAATTGAAACTTGGTACCTCTGAAACAATAACATCCACAT
TCCATCCCCTCCCCAGTCCCTGTTAAACAACCATTTGACTTTATGTCTCTATGAATTTAA
CTACTCTATGTACCTCATATAAATGGAACATATAAGATTTGTTCTTTTGCATCTGGTTTA

6328   GAACCTGGGAGGTGAAGTTTGCAGTGAGCTGAGATTGCTCCACTGCACTCCAGCCTGGCA
ACAGAGCAAGACTCTGTCCAAAAAAAAAACGGCTTGCTTATTTGATTATATAAGATATCTT
TCATAAATTAGATCTCAAATTATACTATTGTTTTGCAGTTTTAGCTTTTATGTTTTAGGG
CAAATCTTAAGTCCTAATTACTTTTTTTTTTATTATTGTGGTAAAATGTATATAACAAAAT
GTACCATTTAATCATTTTAGAATATACGGTTTATGACATTAAGCACATTCACGTTATCAT
[G,A]
CAACCATCACCACTACCCATCCTCAGAACATTTCTCTTCTCGAATTGAAACTTGGTACCT
CTGAAACAATAACATCCACATTCCATCCCCTCCCCAGTCCCTGTTAAACAACCATTTGAC
TTTATGTCTCTATGAATTTAACTACTCTATGTACCTCATATAAATGGAACATATAAGATT
TGTTCTTTTGCATCTGGTTTATTTCATTTAGCATATATTTTTAAGGTTCATCCATGTTGC
AGCATGTGTCAAGATTCTCTTTCTTTTTAAGTCTGAGTCGTATTCCATTGTATGGATATA

6350   AGTGAGCTGAGATTGCTCCACTGCACTCCAGCCTGGCAACAGAGCAAGACTCTGTCCAAA
AAAAAACGGCTTGCTTATTTGATTATATAAGATATCTTTCATAAATTAGATCTCAAATTA

FIGURE 3BBB

TACTATTGTTTTGCAGTTTTAGCTTTTATGTTTTAGGGCAAATCTTAAGTCCTAATTACT
TTTTTTTTATTATTGTGGTAAAATGTATATAACAAAATGTACCATTTAATCATTTTAGAA
TATACGGTTTATGACATTAAGCACATTCACGTTATCATGCAACCATCACCACTACCCATC
[C,T]
TCAGAACATTTCTCTTCTCGAATTGAAACTTGGTACCTCTGAAACAATAACATCCACATT
CCATCCCCTCCCCAGTCCCTGTTAAACAACCATTTGACTTTATGTCTCTATGAATTTAAC
TACTCTATGTACCTCATATAAATGGAACATATAAGATTTGTTCTTTTGCATCTGGTTTAT
TTCATTTAGCATATATTTTTAAGGTTCATCCATGTTGCAGCATGTGTCAAGATTCTCTTT
CTTTTTAAGTCTGAGTCGTATTCCATTGTATGGATATACCACATTTTGTTTATCTTTTCA

6382   CTGGCAACAGAGCAAGACTCTGTCCAAAAAAAAACGGCTTGCTTATTTGATTATATAAGA
        TATCTTTCATAAATTAGATCTCAAATTATACTATTGTTTTGCAGTTTTAGCTTTTATGTT
        TTAGGGCAAATCTTAAGTCCTAATTACTTTTTTTTTTATTATTGTGGTAAAATGTATATAA
        CAAAATGTACCATTTAATCATTTTAGAATATACGGTTTATGACATTAAGCACATTCACGT
        TATCATGCAACCATCACCACTACCCATCCTCAGAACATTTCTCTTCTCGAATTGAAACTT
        [G,T]
        GTACCTCTGAAACAATAACATCCACATTCCATCCCCTCCCCAGTCCCTGTTAAACAACCA
        TTTGACTTTATGTCTCTATGAATTTAACTACTCTATGTACCTCATATAAATGGAACATAT
        AAGATTTGTTCTTTTGCATCTGGTTTATTTCATTTAGCATATATTTTTAAGGTTCATCCA
        TGTTGCAGCATGTGTCAAGATTCTCTTTCTTTTTAAGTCTGAGTCGTATTCCATTGTATG
        GATATACCACATTTTGTTTATCTTTTCATTAGTTGACATTGATTGTCCTCACCTTTTGAT

6434   ATATAAGATATCTTTCATAAATTAGATCTCAAATTATACTATTGTTTTGCAGTTTTAGCT
        TTTATGTTTTAGGGCAAATCTTAAGTCCTAATTACTTTTTTTTTTATTATTGTGGTAAAAT
        GTATATAACAAAATGTACCATTTAATCATTTTAGAATATACGGTTTATGACATTAAGCAC
        ATTCACGTTATCATGCAACCATCACCACTACCCATCCTCAGAACATTTCTCTTCTCGAAT
        TGAAACTTGGTACCTCTGAAACAATAACATCCACATTCCATCCCCTCCCCAGTCCCTGTT
        [A,-]
        AACAACCATTTGACTTTATGTCTCTATGAATTTAACTACTCTATGTACCTCATATAAATG
        GAACATATAAGATTTGTTCTTTTGCATCTGGTTTATTTCATTTAGCATATATTTTTAAGG
        TTCATCCATGTTGCAGCATGTGTCAAGATTCTCTTTCTTTTTAAGTCTGAGTCGTATTCC
        ATTGTATGGATATACCACATTTTGTTTATCTTTTCATTAGTTGACATTGATTGTCCTCAC
        CTTTTGATTTTTGTGAATAAGGCTGCTATAAACATTGGTGTGCAAATATCTGTTCAAGTC

6722   CCAGTCCCTGTTAAACAACCATTTGACTTTATGTCTCTATGAATTTAACTACTCTATGTA
        CCTCATATAAATGGAACATATAAGATTTGTTCTTTTGCATCTGGTTTATTTCATTTAGCA
        TATATTTTTAAGGTTCATCCATGTTGCAGCATGTGTCAAGATTCTCTTTCTTTTTAAGTC
        TGAGTCGTATTCCATTGTATGGATATACCACATTTTGTTTATCTTTTCATTAGTTGACAT
        TGATTGTCCTCACCTTTTGATTTTTGTGAATAAGGCTGCTATAAACATTGGTGTGCAAAT
        [A,G]
        TCTGTTCAAGTCCCTGTTTTCAATTCTTCAGGGTATATACCTAGAAGTGGAAGCACTGGA
        TCATATAATTCCTTGTTTGACTCTCTGAGGAACCATCATACTGTCTTCTACCTAATTATG
        CTTTGTGTTTTAGTAATGGGACACAGCCTGGCATGATGGGCTAGAGTATTGGAAAGGCAT
        GCACAGGTTCAAGTCTCAGCTGTGCCACGTGCCAGTAATCTACATGTTTCTATGAGAAGA
        GTCAAAGAGGATATAGCCTGGTCAACCATTATCAGACACTGGAGTCAGTTTGACTAATTA

6751   TATGTCTCTATGAATTTAACTACTCTATGTACCTCATATAAATGGAACATATAAGATTTG
        TTCTTTTGCATCTGGTTTATTTCATTTAGCATATATTTTTAAGGTTCATCCATGTTGCAG
        CATGTGTCAAGATTCTCTTTCTTTTTAAGTCTGAGTCGTATTCCATTGTATGGATATACC
        ACATTTTGTTTATCTTTTCATTAGTTGACATTGATTGTCCTCACCTTTTGATTTTTGTGA
        ATAAGGCTGCTATAAACATTGGTGTGCAAATATCTGTTCAAGTCCCTGTTTTCAATTCTT
        [C,T]
        AGGGTATATACCTAGAAGTGGAAGCACTGGATCATATAATTCCTTGTTTGACTCTCTGAG
        GAACCATCATACTGTCTTCTACCTAATTATGCTTTGTGTTTTAGTAATGGGACACAGCCT
        GGCATGATGGGCTAGAGTATTGGAAAGGCATGCACAGGTTCAAGTCTCAGCTGTGCCACG
        TGCCAGTAATCTACATGTTTCTATGAGAAGAGTCAAAGAGGATATAGCCTGGTCAACCAT
        TATCAGACACTGGAGTCAGTTTGACTAATTATATGGTGTTCTAAGGAAACTTGAGGTACC

6752   ATGTCTCTATGAATTTAACTACTCTATGTACCTCATATAAATGGAACATATAAGATTTGT
        TCTTTTGCATCTGGTTTATTTCATTTAGCATATATTTTTAAGGTTCATCCATGTTGCAGC
        ATGTGTCAAGATTCTCTTTCTTTTTAAGTCTGAGTCGTATTCCATTGTATGGATATACCA
        CATTTTGTTTATCTTTTCATTAGTTGACATTGATTGTCCTCACCTTTTGATTTTTGTGAA

FIGURE 3CCC

TAAGGCTGCTATAAACATTGGTGTGCAAATATCTGTTCAAGTCCCTGTTTTCAATTCTTC
[A,T]
GGGTATATACCTAGAAGTGGAAGCACTGGATCATATAATTCCTTGTTTGACTCTCTGAGG
AACCATCATACTGTCTTCTACCTAATTATGCTTTGTGTTTTAGTAATGGGACACAGCCTG
GCATGATGGGCTAGAGTATTGGAAAGGCATGCACAGGTTCAAGTCTCAGCTGTGCCACGT
GCCAGTAATCTACATGTTTCTATGAGAAGAGTCAAAGAGGATATAGCCTGGTCAACCATT
ATCAGACACTGGAGTCAGTTTGACTAATTATATGGTGTTCTAAGGAAACTTGAGGTACCA

7070   TGGAAGCACTGGATCATATAATTCCTTGTTTGACTCTCTGAGGAACCATCATACTGTCTT
CTACCTAATTATGCTTTGTGTTTTAGTAATGGGACACAGCCTGGCATGATGGGCTAGAGT
ATTGGAAAGGCATGCACAGGTTCAAGTCTCAGCTGTGCCACGTGCCAGTAATCTACATGT
TTCTATGAGAAGAGTCAAAGAGGATATAGCCTGGTCAACCATTATCAGACACTGGAGTCA
GTTTGACTAATTATATGGTGTTCTAAGGAAACTTGAGGTACCACAAGAAAAGTCTCCAAA
[T,C]
CTAAATAATTACTAATGAATTAATTGAGGGGGAAACTTATTTAACCTTTGTAAGCCTCAG
TTTCTTTGTATGTAAAATGCAGGTAATAATTGGGCATACTTCATTAGGTCTTTGTGAGGA
TTGAATAAATAATGCAAGTAAAACACTTAGCAAAGTATTTCCCATAAAGTAACCACTCAA
TTAATGCTAATTAAGTGTTATTTACTAACATCAGAGTTTCCTAGTGTGAACTCTTTGAAG
TACTTTAAGTTCTGAGAAAAACAAAATTAATTAAATGCAACTCTGTCGATTCCACAGTTA

7306   GTCAGTTTGACTAATTATATGGTGTTCTAAGGAAACTTGAGGTACCACAAGAAAAGTCTC
CAAATCTAAATAATTACTAATGAATTAATTGAGGGGGAAACTTATTTAACCTTTGTAAGC
CTCAGTTTCTTTGTATGTAAAATGCAGGTAATAATTGGGCATACTTCATTAGGTCTTTGT
GAGGATTGAATAAATAATGCAAGTAAAACACTTAGCAAAGTATTTCCCATAAAGTAACCA
CTCAATTAATGCTAATTAAGTGTTATTTACTAACATCAGAGTTTCCTAGTGTGAACTCTT
[T,C]
GAAGTACTTTAAGTTCTGAGAAAAACAAAATTAATTAAATGCAACTCTGTCGATTCCACA
GTTAATTAGACCTATTCATGTTTCTATTGACTGGATTAACAGAACGGCAGATTTTATGGA
TTCTGTTAAAACCTATATAAAAACACTTTAAAAGAAGCCAAGTTATTGACTGCACAAAAA
CATAATCTCATCTGATATCTTTTTTATCCCCCTGAGGTTATTGTGTTTTTGTTTAAGGCA
AAATCAAGAACTAATTGGGATGAAAATAACTAAAGTTTACTTTGTCTGATTTAAGTCCCA

7339   AACTTGAGGTACCACAAGAAAAGTCTCCAAATCTAAATAATTACTAATGAATTAATTGAG
GGGGAAACTTATTTAACCTTTGTAAGCCTCAGTTTCTTTGTATGTAAAATGCAGGTAATA
ATTGGGCATACTTCATTAGGTCTTTGTGAGGATTGAATAAATAATGCAAGTAAAACACTT
AGCAAAGTATTTCCCATAAAGTAACCACTCAATTAATGCTAATTAAGTGTTATTTACTAA
CATCAGAGTTTCCTAGTGTGAACTCTTTGAAGTACTTTAAGTTCTGAGAAAAACAAAATT
[A,G]
ATTAAATGCAACTCTGTCGATTCCACAGTTAATTAGACCTATTCATGTTTCTATTGACTG
GATTAACAGAACGGCAGATTTTATGGATTCTGTTAAAACCTATATAAAAACACTTTAAAA
GAAGCCAAGTTATTGACTGCACAAAAACATAATCTCATCTGATATCTTTTTTATCCCCCT
GAGGTTATTGTGTTTTTGTTTAAGGCAAAATCAAGAACTAATTGGGATGAAAATAACTAA
AGTTTACTTTGTCTGATTTAAGTCCCAAACTGACTAATAAGTAATCCCATTTGATCAACA

7531   CCCATAAAGTAACCACTCAATTAATGCTAATTAAGTGTTATTTACTAACATCAGAGTTTC
CTAGTGTGAACTCTTTGAAGTACTTTAAGTTCTGAGAAAAACAAAATTAATTAAATGCAA
CTCTGTCGATTCCACAGTTAATTAGACCTATTCATGTTTCTATTGACTGGATTAACAGAA
CGGCAGATTTTATGGATTCTGTTAAAACCTATATAAAAACACTTTAAAAGAAGCCAAGTT
ATTGACTGCACAAAAACATAATCTCATCTGATATCTTTTTTATCCCCCTGAGGTTATTGT
[G,A]
TTTTTGTTTAAGGCAAAATCAAGAACTAATTGGGATGAAAATAACTAAAGTTTACTTTGT
CTGATTTAAGTCCCAAACTGACTAATAAGTAATCCCATTTGATCAACAGATTCAGTGAAA
ACTGTCCCCCATTCTCAACTACCATATGGATATTCTGAGAAATAATTAATGATGCAGAAA
AACATTTTTTGTTTTCTGAAATAAAAGAATAGACGTGCAAGTGACACTTCTTTTTAATGC
TTACAACCTTTTTTTAAAAATCTACTTTATTTTCTCTATCTGAATGCACTAGATTTTGTT

8902   AAGTCCAATTTATGGCAAGGGTTTTAATTTGTAAGGGCTTTATTTCTCCATACAAAGGGA
TTGGAGAAACAAACTAGAAAGCCAGAAAACAGACCACAAACACTGAGCTAGTGGTTCCAA
CTGGAGTGTTCCCTGAGCAGTGACTTATGAATACTTGTTTAGAAGAATCAACTCAAACAA
ATTTAGGAAAGTCACATCCTGCCTTTAGAGCTTCCAGTGTTTGTTAGCATATTAAAGTCT
CTGAAATGACCTACAATATTGAAATCTCAGTCTTCTGCTATTTTTAATATTTATTTCAAA
[A,G]

FIGURE 3DDD

TGAAATAATTTTTGTGAAAAACATTTTAATGTCTGTGGCTCATAATATTCTGTGGATCTC
AGTTTGGGAAATGAAAGATTATAATCGTATCTACTCTTTATCTGTTGGAAACATCTTTCC
ATTTATTTTTCCTGCTGGTTTAATGGCAACAAATTTTTACATGTGAAATATTTGTAATGT
GATTTATATGAAAAAATGTAATTTTCTTATTACACGATCAAAAGTGGTTATGCTCCTCTG
TAAGTTTTTCCTTACAAGTTTTTATGTTGCATAATTTATATCTATTTGGTTTAATGAGTA

9471   GCATAATTTATATCTATTTGGTTTAATGAGTACAACACAAGATAGCTCAGTTTAATTCTG
          GGATGTTGGATGTTTCTAGTTAAAGTACAAGTTGGATTTGATGAAAATTCATTGCTTCTT
          TATGATTTTTTTAAAACTCAAGAACATGTTAGTTAAAGAGTGTCTTCTGAACAAATTCTTG
          TGAAGTAGTTGCTGATTATTAAGTAACACTCATGCTACCGTAACTTTTTATACTATCCAA
          AGCTATAGACATTTTTAATTTTTCAACTTGCAACTACCTAGGTTGAAAAAATTAAATCTGCA
          [G,A]
          GCCAGTTTCATTATTCAGACAATTTGGTTATCACTTCAAGCCTACTATCTTCAAAGAAAA
          TGGGAGTGCAGGCCTTCATGGGAGCTGACTTCTGCTGTATGGCCTTGCAAATGTCAACTC
          GATTAGAGTGACCAGTGTTAGCCCTCAATTCACAAACTCAGGTCCCATGAAATATACACG
          GATTTCTACTATGCATTACTATGTGACCATTCATGGAAGTTTCGTTTGGAAACACAGACA
          TTAAAAAGCCAGTCATGGAATAACATTCTTGTTAAAACAGGACATTGGCAAAAAGGACTA

10023   GTCATGGAATAACATTCTTGTTAAAACAGGACATTGGCAAAAAGGACTAGAAAACTTCTG
          GCTATAGATTTTGAATCCAATAGCCTTGCATAGGCTTTTCTGTTTCCTCCTAAACTATGT
          CTTCTGTCCTTTCTGGAGGCATATTTATAGTAAAATAAACAAAATTAACCTTGTTTTACA
          CTTGAGTAACCTATACCTTTGGTTATTTACGAGAATTACTTAAAGCAGAGTTGGCAACTT
          TTTCTGTGATGGGCCTGATACTAAATATTTTACACTTTCCAAGTAATACAGTCTCTGTCA
          [C,T]
          AACTACTCAACTCTGCCACTGTAGCATAAAAGCACACTTAGACAATGCAGAAACAAATGA
          ACATGGCTTTGTTCCAATAAAACTTTATTTATGGACACTGAAATGTGAATTTCAAAAATA
          TTTTTTGCATAAGATCAAATATTATTCTTTTGATTTTTTTTCCAATCAATAAAAAGTGTAA
          AAATTGGCCGGGCATGGTGGCTCATGCCTGTAATCCCAGCACTTTGGGAGGCCGAGGTGG
          GCAGATCACCTGAGGTCACGAGTTCGAGACCAGCCTGACCAACATGGAGAAACCCTGTCT

10594   CAGCCTGACCAACATGGAGAAACCCTGTCTATATTAAAAAATACAAAATTAGCTGGGTGTG
          GTGGGGCCTACCTGTAATCCCAGCTACTCGGGAGGCTGAGGCAGGAGAATCGCTTGAATG
          CAGGAGGCAGAGGTTGCGGTGAGCCCAGATTGCACCATTGCACTCCAGCCGGGGCAACAA
          GAGCAAAACTCCGTCTCAAAAAAAAAAAAAAAAAAAAAGTGTAAAAACCATTCTTAGTTCA
          TGAGCTATACAAAAATAGATAGTGAGTTAGATTTGGCCCATGGGGCTTATTTTGCTGACT
          [C,G]
          CTGCTCTAAGCATCTTGCAGACATTTCTTCATATGCCCTAGGAGATTTCTGATATCCCCT
          CATAATACCCTGGCCTTACACCAAGACTACAATCTGTTCTTTGCAGATGCTTAATAAATT
          CATTCTTCCCTGTCATTCAGTTGATCTGTGTGAGCCAGTGGAAATACTTGGGCCAATAAA
          TCTAGTGTGTTTGAGGGTAAAATATGCTATTTTTGTAAGATATATTATTTAATGGCCACA
          CAACCTAAATTCAATTAAATGGTTACAACCTGTAACGCATTTAAAAATATGACTAGGCAGA

11233   CATCCAACAGTTGATGTTGATCCCCCCATCCTGCCCCACTGTTCTACTTTGCAATTTGTT
          TGAAAGAAATTGTCAATATATTTCTGACTTCTGAGCAAATCCATGAATCGGGATCCAGCA
          ACAGGAAAAGAAGCTGTTGCTGCCCATTGCTTGGTTTTGGCACCAGGAATGGATAAATCC
          CAGACTTCCTGGGGCACGTGTTTTATAAAAGGGAAGTGCTGACAGTGCAAACAGCTGCCA
          TCAATTGGCCTTGGAGACTACTTCCCTGGAGAAGCTCCAATTATATTCTTAAAGGACCCA
          [C,T]
          CAAGCTCTTCAAGTGTTAGTGGCAACCATTTGCTGCCAACCATTTGAAATGATGAAGTAA
          TTTTTTTTTATTAGTGGATCCTAAGTGATAGGCTCTAGAACTGATCTTCAACCTTAACTA
          ATATCATGGCATCAGAGGGCTACAGATTAAATCAGTGGTTCCCAGTCACTCTCTGTGGAC
          AAGTAGCAACTACGACAAAGCTTTTCTTAGTCTATGGTGGAAGAGAAAAATTAGGACAAT
          GTAATAAGCATCCCATAAACTTATTAAACCTATTAAAATTTAATTTTAAGATTATGTCAT

11295   AAAGAAATTGTCAATATATTTCTGACTTCTGAGCAAATCCATGAATCGGGATCCAGCAAC
          AGGAAAAGAAGCTGTTGCTGCCCATTGCTTGGTTTTGGCACCAGGAATGGATAAATCCCA
          GACTTCCTGGGGCACGTGTTTTATAAAAGGGAAGTGCTGACAGTGCAAACAGCTGCCATC
          AATTGGCCTTGGAGACTACTTCCCTGGAGAAGCTCCAATTATATTCTTAAAGGACCCACC
          AAGCTCTTCAAGTGTTAGTGGCAACCATTTGCTGCCAACCATTTGAAATGATGAAGTAAT
          [-,A,T]
          TTTTTTTTATTAGTGGATCCTAAGTGATAGGCTCTAGAACTGATCTTCAACCTTAACTAAT
          ATCATGGCATCAGAGGGCTACAGATTAAATCAGTGGTTCCCAGTCACTCTCTGTGGACAA

FIGURE 3EEE

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          GTAGCAACTACGACAAAGCTTTTCTTAGTCTATGGTGGAAGAGAAAAATTAGGACAATGT
          AATAAGCATCCCATAAACTTATTAAACCTATTAAAATTTAATTTTAAGATTATGTCATTT
          TTTGTATGTGTGTATGCTTAGTATTTATGGATTGTGGAAATAGAATTTTTTTTTTTATAGT
```

11534   CAAGCTCTTCAAGTGTTAGTGGCAACCATTTGCTGCCAACCATTTGAAATGATGAAGTAA
          TTTTTTTTTTATTAGTGGATCCTAAGTGATAGGCTCTAGAACTGATCTTCAACCTTAACTA
          ATATCATGGCATCAGAGGGCTACAGATTAAATCAGTGGTTCCCAGTCACTCTCTGTGGAC
          AAGTAGCAACTACGACAAAGCTTTTCTTAGTCTATGGTGGAAGAGAAAAATTAGGACAAT
          GTAATAAGCATCCCATAAACTTATTAAACCTATTAAAATTTAATTTTAAGATTATGTCAT
          [-,T]
          TTTGTATGTGTGTATGCTTAGTATTTATGGATTGTGGAAATAGAATTTTTTTTTTATAG
          TGAGAACCTAGGTAAGTGACTTACCTCTCTGATCCCCCATTTTCTCATATGTAGAAGGGG
          GCTAATAATAGTATCTGTCTCATAGTTTTTGTGAGAATAAAAAAAATTGTCCAGGTAAAAT
          GCTTAGCTGGTGACTGGCACACAGTAATTGCTCAATAAATGTTAGCTATTATTGCTATCA
          TTATATAATCATCATGGTTTCCAATGCCTTTACTTGGCAAATAAAAGAACAAAAGTCACC

11757   AGAAAAATTAGGACAATGTAATAAGCATCCCATAAACTTATTAAACCTATTAAAATTTAA
          TTTTAAGATTATGTCATTTTTTGTATGTGTGTATGCTTAGTATTTATGGATTGTGGAAAT
          AGAATTTTTTTTTTTATAGTGAGAACCTAGGTAAGTGACTTACCTCTCTGATCCCCCATTT
          TCTCATATGTAGAAGGGGGCTAATAATAGTATCTGTCTCATAGTTTTTGTGAGAATAAAA
          AAATTGTCCAGGTAAAATGCTTAGCTGGTGACTGGCACACAGTAATTGCTCAATAAATGT
          [T,C]
          AGCTATTATTGCTATCATTATATAATCATCATGGTTTCCAATGCCTTTACTTGGCAAATA
          AAAGAACAAAAGTCACCCGATATTGATCTCCCTTTTCTTCCCTAGTTTTCTGGGGGGTGG
          GAGGCAGAGACCGAATTTTCTGATCTGTGAAATCTGAATTTATCATTGTAATTTTCCATA
          AGTGCTATGTAGAGAACTCATTTAAGTTGCTGGGATGAAAAAAAAATCAAAAGTGGCCTAT
          TGTGCTGGGTGCAGTGGTTCACGCCTGCAATCCCAGCACTTTGGGAGGCTGAGGGGGGTG

11951   GGGGGCTAATAATAGTATCTGTCTCATAGTTTTTGTGAGAATAAAAAAAATTGTCCAGGTA
          AAATGCTTAGCTGGTGACTGGCACACAGTAATTGCTCAATAAATGTTAGCTATTATTGCT
          ATCATTATATAATCATCATGGTTTCCAATGCCTTTACTTGGCAAATAAAAGAACAAAAGT
          CACCCGATATTGATCTCCCTTTTCTTCCCTAGTTTTCTGGGGGGTGGGAGGCAGAGACCG
          AATTTTCTGATCTGTGAAATCTGAATTTATCATTGTAATTTTCCATAAGTGCTATGTAGA
          [G,A]
          AACTCATTTAAGTTGCTGGGATGAAAAAAAAATCAAAAGTGGCCTATTGTGCTGGGTGCAG
          TGGTTCACGCCTGCAATCCCAGCACTTTGGGAGGCTGAGGGGGGTGGATCGCCTGAGGTC
          AGGAGTTCAAGACCAGCCTGGCCAACATGGTGAAACCTTGACTCTACTAAAAATACAAAA
          ATTAGCCTGGCATGATGGTGGGCACCTGTAATCCCAGCTACTCAGGAGGCTGAGGCAGGA
          GAATCCCTTGAACCCAGGAGGTGGAGGATTCAGTGAGCCGAGATCTACTGCACTCCAGCC

12901   AGTGATTATGGGACAGCTAAATGATGGTGCCAAGTAGGAGCTGGAGTAGAGTATCCAGCA
          ATGAGTGGAAACATCTGGGATGGAGACAGAAAGACACGGGTATTAATTCTACGGGGATGG
          CTAAGTCTGCTCTGAGAGACAGTGTGGAGACCAAGGAGAAGAGGAATCCTAATATTTAGA
          AACAAGGCAGTGGATAGCAATCTAGCTATGGAAAGTGGAAGGAAAGAGATAGTTGATCAT
          CCAGTTCAACACTACTCTTGTTGTAGTTCACTTATGTTGAATGCTTCTGTGTGACTAAGT
          [C,A]
          GGTGAGAAAAATCTATGGGAGTAGGCAACATGGAGGATGTTGGTATTCACAAAAGCAGTT
          TAGTGGAGTGTGGAGGCCTGAGCCAGACTAGAATGAGTTAGGAGTAGATGGAAGATAAGA
          ATGCAGATATGGGCCCAGCGCGGTGGCTCACGCCTGTAATCCCAGCACTTTGGGAGGCCA
          AGGTGAGCAGATCACAAGGTCAGGAGATCGAGACCATCCTGGCTAACACCGTGAAACCCC
          ATCTCTACTAAAAATACAAAAAATTAGCCGGGCCTGGTGGCGGGTGCCTGTAGTCCCAGC

13040   CAGTGTGGAGACCAAGGAGAAGAGGAATCCTAATATTTAGAAACAAGGCAGTGGATAGCA
          ATCTAGCTATGGAAAGTGGAAGGAAAGAGATAGTTGATCATCCAGTTCAACACTACTCTT
          GTTGTAGTTCACTTATGTTGAATGCTTCTGTGTGACTAAGTCGGTGAGAAAAATCTATGG
          GAGTAGGCAACATGGAGGATGTTGGTATTCACAAAAGCAGTTTAGTGGAGTGTGGAGGCC
          TGAGCCAGACTAGAATGAGTTAGGAGTAGATGGAAGATAAGAATGCAGATATGGGCCCAG
          [C,T]
          GCGGTGGCTCACGCCTGTAATCCCAGCACTTTGGGAGGCCAAGGTGAGCAGATCACAAGG
          TCAGGAGATCGAGACCATCCTGGCTAACACCGTGAAACCCCATCTCTACTAAAAATACAA
          AAAATTAGCCGGGCCTGGTGGCGGGTGCCTGTAGTCCCAGCTACTCGGGAGGCTGAGGCA
          GGAGAATGGCGTGAACCCGGGAGGTGGAGCTGGCAGTGAGCCGAGATGGTGCCACTGCAC
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# FIGURE 3FFF

TCCAGCCTGGGCAACAGAGCAAGACTCCATCTCAAAAAAAAAAAAAAAAAAAAAGAATGCAG

13081    AACAAGGCAGTGGATAGCAATCTAGCTATGGAAAGTGGAAGGAAAGAGATAGTTGATCAT
CCAGTTCAACACTACTCTTGTTGTAGTTCACTTATGTTGAATGCTTCTGTGTGACTAAGT
CGGTGAGAAAAATCTATGGGAGTAGGCAACATGGAGGATGTTGGTATTCACAAAAGCAGT
TTAGTGGAGTGTGGAGGCCTGAGCCAGACTAGAATGAGTTAGGAGTAGATGGAAGATAAG
AATGCAGATATGGGCCCAGCGCGGTGGCTCACGCCTGTAATCCCAGCACTTTGGGAGGCC
[A,G]
AGGTGAGCAGATCACAAGGTCAGGAGATCGAGACCATCCTGGCTAACACCGTGAAACCCC
ATCTCTACTAAAAATACAAAAAATTAGCCGGGCCTGGTGGCGGGTGCCTGTAGTCCCAGC
TACTCGGGAGGCTGAGGCAGGAGAATGGCGTGAACCCGGGAGGTGGAGCTGGCAGTGAGC
CGAGATGGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTCCATCTCAAAAAAAA
AAAAAAAAAAAGAATGCAGATATGGCAAGTATAGACAAGCTTCAAGAAGTTTGGTCTAAA

13173    TATGTTGAATGCTTCTGTGTGACTAAGTCGGTGAGAAAAATCTATGGGAGTAGGCAACAT
GGAGGATGTTGGTATTCACAAAAGCAGTTTAGTGGAGTGTGGAGGCCTGAGCCAGACTAG
AATGAGTTAGGAGTAGATGGAAGATAAGAATGCAGATATGGGCCCAGCGCGGTGGCTCAC
GCCTGTAATCCCAGCACTTTGGGAGGCCAAGGTGAGCAGATCACAAGGTCAGGAGATCGA
GACCATCCTGGCTAACACCGTGAAACCCCATCTCTACTAAAAATACAAAAAATTAGCCGG
[G,T]
CCTGGTGGCGGGTGCCTGTAGTCCCAGCTACTCGGGAGGCTGAGGCAGGAGAATGGCGTG
AACCCGGGAGGTGGAGCTGGCAGTGAGCCGAGATGGTGCCACTGCACTCCAGCCTGGGCA
ACAGAGCAAGACTCCATCTCAAAAAAAAAAAAAAAAAAAAAGAATGCAGATATGGCAAGTAT
AGACAAGCTTCAAGAAGTTTGGTCTAAAAGGAAGCGGAGAAATAAACAAAGAGATGATGC
CTAATATAATTCAGCTAAATGTAATATAATGGATTTTTTTAAGATGAGGTACTAGAGCAT

13272    TGGAGGCCTGAGCCAGACTAGAATGAGTTAGGAGTAGATGGAAGATAAGAATGCAGATAT
GGGCCCAGCGCGGTGGCTCACGCCTGTAATCCCAGCACTTTGGGAGGCCAAGGTGAGCAG
ATCACAAGGTCAGGAGATCGAGACCATCCTGGCTAACACCGTGAAACCCCATCTCTACTA
AAAATACAAAAAATTAGCCGGGCCTGGTGGCGGGTGCCTGTAGTCCCAGCTACTCGGGAG
GCTGAGGCAGGAGAATGGCGTGAACCCGGGAGGTGGAGCTGGCAGTGAGCCGAGATGGTG
[C,T]
CACTGCACTCCAGCCTGGGCAACAGAGCAAGACTCCATCTCAAAAAAAAAAAAAAAAAAAA
GAATGCAGATATGGCAAGTATAGACAAGCTTCAAGAAGTTTGGTCTAAAAGGAAGCGGAG
AAATAAACAAAGAGATGATGCCTAATATAATTCAGCTAAATGTAATATAATGGATTTTTT
TAAGATGAGGTACTAGAGCATGTAATATAAATCTATTAAATTGGGTGGCCAGGAACCAGG
ACTGGCTCATCAGCATGGACCAGGCTAGACGCACAGGGCCTTATATCCAGAAGGACATCA

13333    GGCCCAGCGCGGTGGCTCACGCCTGTAATCCCAGCACTTTGGGAGGCCAAGGTGAGCAGA
TCACAAGGTCAGGAGATCGAGACCATCCTGGCTAACACCGTGAAACCCCATCTCTACTAA
AAATACAAAAAATTAGCCGGGCCTGGTGGCGGGTGCCTGTAGTCCCAGCTACTCGGGAGG
CTGAGGCAGGAGAATGGCGTGAACCCGGGAGGTGGAGCTGGCAGTGAGCCGAGATGGTGC
CACTGCACTCCAGCCTGGGCAACAGAGCAAGACTCCATCTCAAAAAAAAAAAAAAAAAAAA
[A,-,G]
AATGCAGATATGGCAAGTATAGACAAGCTTCAAGAAGTTTGGTCTAAAAGGAAGCGGAGA
AATAAACAAAGAGATGATGCCTAATATAATTCAGCTAAATGTAATATAATGGATTTTTTT
AAGATGAGGTACTAGAGCATGTAATATAAATCTATTAAATTGGGTGGCCAGGAACCAGGA
CTGGCTCATCAGCATGGACCAGGCTAGACGCACAGGGCCTTATATCCAGAAGGACATCAC
CTTTGGGTTTTAATGCTCTGCACTTGCTGTCTCCAAATTCTAACTGTCTCTTAGGCTCTC

13485    GTGCCTGTAGTCCCAGCTACTCGGGAGGCTGAGGCAGGAGAATGGCGTGAACCCGGGAGG
TGGAGCTGGCAGTGAGCCGAGATGGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGA
CTCCATCTCAAAAAAAAAAAAAAAAAAAAAGAATGCAGATATGGCAAGTATAGACAAGCTTC
AAGAAGTTTGGTCTAAAAGGAAGCGGAGAAATAAACAAAGAGATGATGCCTAATATAATT
CAGCTAAATGTAATATAATGGATTTTTTTTAAGATGAGGTACTAGAGCATGTAATATAAAT
[C,A]
TATTAAATTGGGTGGCCAGGAACCAGGACTGGCTCATCAGCATGGACCAGGCTAGACGCA
CAGGGCCTTATATCCAGAAGGACATCACCTTTGGGTTTTAATGCTCTGCACTTGCTGTCT
CCAAATTCTAACTGTCTCTTAGGCTCTCATCAACACCCACCTCCATATCCAGATATTGAG
TACCTCAGGGAGTTCAATTTGGAAGCAAATGATGTGAAAATGTACTTTACTATCCAGTAA
CATTCTTGTTAGGGAGTGTTGGCAGAGATTGTCGAACAACCATAATGCATTTTATCATTC

# FIGURE 3GGG

13933    CATCAACACCCACCTCCATATCCAGATATTGAGTACCTCAGGGAGTTCAATTTGGAAGCA
AATGATGTGAAAATGTACTTTACTATCCAGTAACATTCTTGTTAGGGAGTGTTGGCAGAG
ATTGTCGAACAACCATAATGCATTTTATCATTCGATCAGTCTACAATTTAAACATAGCAG
GACTGGACAGAGGCACAGGAAGATTAAGCCACTGACCTTAAGTCAGACAGTCACATGGGT
AGATCCGGAATCTTGATCTAAAATGAATACCATTTTTTTCAGTTATAGCTATCTTCCCAGG
[A,T]
TGGCCAACCAGAATGCATATATAAAATTTCAAAAACAAACATTGGGAATTGCTCTTCAGC
AAGAATACATCAAACACCCATTATGTGCCTAACTCTAAATCTTACTTTCAGAGAGCTAAA
AACAATTTCATTTCACAGTGACATTCATCTTCGCTTCTGCCGTAACTCACATGCATATGC
CTTAGACCACATTATTAATGAAGTATTGGGGGGTTCCATCTAGAGCACCTTTTCTTCCCT
GGAGTTAATCATCCAGTTCAGCACCACTCTTGAGCTTTGCTTAGCTTCTTCTACCCATTT

14086    GATCAGTCTACAATTTAAACATAGCAGGACTGGACAGAGGCACAGGAAGATTAAGCCACT
GACCTTAAGTCAGACAGTCACATGGGTAGATCCGGAATCTTGATCTAAAATGAATACCAT
TTTTTCAGTTATAGCTATCTTCCCAGGATGGCCAACCAGAATGCATATATAAAATTTCAA
AAACAAACATTGGGAATTGCTCTTCAGCAAGAATACATCAAACACCCATTATGTGCCTAA
CTCTAAATCTTACTTTCAGAGAGCTAAAAACAATTTCATTTCACAGTGACATTCATCTTC
[G,A]
CTTCTGCCGTAACTCACATGCATATGCCTTAGACCACATTATTAATGAAGTATTGGGGGG
TTCCATCTAGAGCACCTTTTCTTCCCTGGAGTTAATCATCCAGTTCAGCACCACTCTTGA
GCTTTGCTTAGCTTCTTCTACCCATTTGGATTTTAAGGACAACAATTCCAATGGCCTTTA
TCCATGTATTTAACAATTCATTATGAGCCAGGTGAAGTGGATCACACCTCTAATCCCAAC
ACTTTGGGAGGCTGAGGCAGGTGGATCGCTGGAGCCCAGGAGTTCACAACCAGCCTGGGC

14094    TACAATTTAAACATAGCAGGACTGGACAGAGGCACAGGAAGATTAAGCCACTGACCTTAA
GTCAGACAGTCACATGGGTAGATCCGGAATCTTGATCTAAAATGAATACCATTTTTTTCAG
TTATAGCTATCTTCCCAGGATGGCCAACCAGAATGCATATATAAAATTTCAAAAACAAAC
ATTGGGAATTGCTCTTCAGCAAGAATACATCAAACACCCATTATGTGCCTAACTCTAAAT
CTTACTTTCAGAGAGCTAAAAACAATTTCATTTCACAGTGACATTCATCTTCGCTTCTGC
[C,T]
GTAACTCACATGCATATGCCTTAGACCACATTATTAATGAAGTATTGGGGGGTTCCATCT
AGAGCACCTTTTCTTCCCTGGAGTTAATCATCCAGTTCAGCACCACTCTTGAGCTTTGCT
TAGCTTCTTCTACCCATTTGGATTTTAAGGACAACAATTCCAATGGCCTTTATCCATGTA
TTTAACAATTCATTATGAGCCAGGTGAAGTGGATCACACCTCTAATCCCAACACTTTGGG
AGGCTGAGGCAGGTGGATCGCTGGAGCCCAGGAGTTCACAACCAGCCTGGGCAACATGGT

14141    CCACTGACCTTAAGTCAGACAGTCACATGGGTAGATCCGGAATCTTGATCTAAAATGAAT
ACCATTTTTTTCAGTTATAGCTATCTTCCCAGGATGGCCAACCAGAATGCATATATAAAAT
TTCAAAAACAAACATTGGGAATTGCTCTTCAGCAAGAATACATCAAACACCCATTATGTG
CCTAACTCTAAATCTTACTTTCAGAGAGCTAAAAACAATTTCATTTCACAGTGACATTCA
TCTTCGCTTCTGCCGTAACTCACATGCATATGCCTTAGACCACATTATTAATGAAGTATT
[G,-]
GGGGGTTCCATCTAGAGCACCTTTTCTTCCCTGGAGTTAATCATCCAGTTCAGCACCACT
CTTGAGCTTTGCTTAGCTTCTTCTACCCATTTGGATTTTAAGGACAACAATTCCAATGGC
CTTTATCCATGTATTTAACAATTCATTATGAGCCAGGTGAAGTGGATCACACCTCTAATC
CCAACACTTTGGGAGGCTGAGGCAGGTGGATCGCTGGAGCCCAGGAGTTCACAACCAGCC
TGGGCAACATGGTGAGACTCCATCTCTACCATTTTTTTTTTTAATTAGTTGGGTATGGTGG

14831    ACAATTTATTTCATCATCATTGTCATCATCATTGTCACTGCTCACTCTTCAACATTTTTT
AGGTCAACTTAATTAATATGATACCTTGTGGGATAATTTTTATTTATTTTTATAAAATAT
TGAAGTTTTTGCCACTTTGATAACTTCTTCATTTTCTGTCCAGAGTATAACATACCAGGG
AAAAGGCTCTAAAATAAGGCTTGAGGTATTAAAAAGATCTTCTGTTTAAGTCTTATGTTC
CTAATCAATAACTAGAATTGGCCTGATTGCTTTCCTCAGTGGGTTTTCTGGTAGTCCTGA
[T,C]
ATGATATCGAGGCTGTCATATAGTCCTGAAATATCCTATCATTAACATTTGTGGTGGTAT
CTGATATAAAGGTAGATGAACTTCATTGCAGCTATTCTTAGGAAATGCGTATTTAAATGC
ATAGTTAAAAGCAAGATTTACAATTATAGAAGGAATGCAAATGAGTTGTAGAAAGCTCAT
AAAATAAAAATCAAGAAGAAAGAATTACCCATCATGCCTCAGCCCAGTGATAACCACTGC
TAATATTTTTGGCTGTTTTTCATTTGCAACCCCATCTCCATTCTAGCAGCCCTCATCCCTC

15319    AAATCAAGAAGAAAGAATTACCCATCATGCCTCAGCCCAGTGATAACCACTGCTAATATT
TTTGGCTGTTTTTCATTTGCAACCCCATCTCCATTCTAGCAGCCCTCATCCCTCCTACCCA

# FIGURE 3HHH

CTATGTTTTTCACTATATTTCTTGTTTAAATTTACTTAATTATTTGTTAATTATGTTTTT
CCTCTCACTAGAAAGTGAACTCCATGAGGGCCAGGGATTTTTGCTATTTTGTTCACTTTT
GTATCCTTAGCACCTACTTTGTTGATTAAGTGAATGCATTAATGATCTATTTTTAATCTG
[T,C]
GTATGTGTATAAAAGACACTTGATATATCTGGGATGATATTCAATATACTTTTGTATCCT
CATTTTTCACCATAGGTAGTTTATGTCAATTCCTTGAAATTTGTTGATTTTCTTGAATAAT
TTAGCAGTTGTACAATTCTAAAACATAAATATAATTTGCTTAAATATACATACCATTTTA
AACATATTTAAATGTGAAAATACAGTTGAGTTCTCTTAGATTGCAATTTTGTAACTTTTG
ATAATCCTTTGATCCTGAAAAAAATTTTTTGGCATGAGGGAAGAGATGAATATTTCTTTT

15321    ATCAAGAAGAAAGAATTACCCATCATGCCTCAGCCCAGTGATAACCACTGCTAATATTTT
TGGCTGTTTTCATTTGCAACCCCATCTCCATTCTAGCAGCCCTCATCCCTCCTACCCACT
ATGTTTTTCACTATATTTCTTGTTTAAATTTACTTAATTATTTGTTAATTATGTTTTTCC
TCTCACTAGAAAGTGAACTCCATGAGGGCCAGGGATTTTTGCTATTTTGTTCACTTTTGT
ATCCTTAGCACCTACTTTGTTGATTAAGTGAATGCATTAATGATCTATTTTTAATCTGTG
[T,C]
ATGTGTATAAAAGACACTTGATATATCTGGGATGATATTCAATATACTTTTGTATCCTCA
TTTTCACCATAGGTAGTTTATGTCAATTCCTTGAAATTTGTTGATTTTCTTGAATAATTT
AGCAGTTGTACAATTCTAAAACATAAATATAATTTGCTTAAATATACATACCATTTTAAA
CATATTTAAATGTGAAAATACAGTTGAGTTCTCTTAGATTGCAATTTTGTAACTTTTGAT
AATCCTTTGATCCTGAAAAAAATTTTTTGGCATGAGGGAAGAGATGAATATTTCTTTTGG

15335    ATTACCCATCATGCCTCAGCCCAGTGATAACCACTGCTAATATTTTTGGCTGTTTTCATT
TGCAACCCCATCTCCATTCTAGCAGCCCTCATCCCTCCTACCCACTATGTTTTTCACTAT
ATTTCTTGTTTAAATTTACTTAATTATTTGTTAATTATGTTTTTCCTCTCACTAGAAAGT
GAACTCCATGAGGGCCAGGGATTTTTGCTATTTTGTTCACTTTTGTATCCTTAGCACCTA
CTTTGTTGATTAAGTGAATGCATTAATGATCTATTTTTAATCTGTGTATGTGTATAAAAG
[A,G]
CACTTGATATATCTGGGATGATATTCAATATACTTTTGTATCCTCATTTTCACCATAGGT
AGTTTATGTCAATTCCTTGAAATTTGTTGATTTTCTTGAATAATTTAGCAGTTGTACAAT
TCTAAAACATAAATATAATTTGCTTAAATATACATACCATTTTAAACATATTTAAATGTG
AAAATACAGTTGAGTTCTCTTAGATTGCAATTTTGTAACTTTTGATAATCCTTTGATCCT
GAAAAAAATTTTTTGGCATGAGGGAAGAGATGAATATTTCTTTTGGAGTATTTAAATCAT

15477    ATTATTTGTTAATTATGTTTTTTCCTCTCACTAGAAAGTGAACTCCATGAGGGCCAGGGAT
TTTTGCTATTTTGTTCACTTTTGTATCCTTAGCACCTACTTTGTTGATTAAGTGAATGCA
TTAATGATCTATTTTTAATCTGTGTATGTGTATAAAAGACACTTGATATATCTGGGATGA
TATTCAATATACTTTTGTATCCTCATTTTTCACCATAGGTAGTTTATGTCAATTCCTTGAA
ATTTGTTGATTTTCTTGAATAATTTAGCAGTTGTACAATTCTAAAACATAAATATAATTT
[G,A]
CTTAAATATACATACCATTTTAAACATATTTAAATGTGAAAATACAGTTGAGTTCTCTTA
GATTGCAATTTTGTAACTTTTGATAATCCTTTGATCCTGAAAAAAATTTTTTGGCATGAG
GGAAGAGATGAATATTTCTTTTGGAGTATTTAAATCATCTCTGCAATAATCCTTTGATCC
TGAAAAAAAATTTGTGGCATGAGGGAAGAGAAGAATATTTCTTTTGGAGTGTTTAAATCA
TCTCTACAATTAATAATATCTAAAGCAGTTTGGTTGGTTTATTTAGGTAGGATTAATTTT

15650    GGGATGATATTCAATATACTTTTGTATCCTCATTTTCACCATAGGTAGTTTATGTCAATT
CCTTGAAATTTGTTGATTTTCTTGAATAATTTAGCAGTTGTACAATTCTAAAACATAAAT
ATAATTTGCTTAAATATACATACCATTTTAAACATATTTAAATGTGAAAATACAGTTGAG
TTCTCTTAGATTGCAATTTTGTAACTTTTGATAATCCTTTGATCCTGAAAAAAATTTTTT
GGCATGAGGGAAGAGATGAATATTTCTTTTGGAGTATTTAAATCATCTCTGCAATAATCC
[T,C]
TTGATCCTGAAAAAAAATTTGTGGCATGAGGGAAGAGAAGAATATTTCTTTTGGAGTGTT
TAAATCATCTCTACAATTAATAATATCTAAAGCAGTTTGGTTGGTTTATTTAGGTAGGAT
TAATTTTCAGTATGAATATTATTTAAAAAACAAATATAGTCAGTTGAATTGCTGTGGAGG
TTTCTGTACGATTTACTCAAAGCTGGCTCTTTTTCTGTACGCACTACCACGCCCGGCTAA
TTTTTGCATTTTTTTTGGTAGAGATGGGGGTTTCACCATGTTGGCCAGGCTGGTCTTGAAC

15880    AAAATTTTTTGGCATGAGGGAAGAGATGAATATTTCTTTTGGAGTATTTAAATCATCTCT
GCAATAATCCTTTGATCCTGAAAAAAAATTTGTGGCATGAGGGAAGAGAAGAATATTTCT
TTTGGAGTGTTTAAATCATCTCTACAATTAATAATATCTAAAGCAGTTTGGTTGGTTTAT
TTAGGTAGGATTAATTTTCAGTATGAATATTATTTAAAAAACAAATATAGTCAGTTGAAT

# FIGURE 3III

TGCTGTGGAGGTTTCTGTACGATTTACTCAAAGCTGGCTCTTTTTCTGTACGCACTACCA
[C,T]
GCCCGGCTAATTTTTTGCATTTTTTTTGGTAGAGATGGGGGTTTCACCATGTTGGCCAGGCT
GGTCTTGAACTCCTGATCTCAAGTGATCCACCCACCTCAGCCTCTCAAGGTGCTGGGATT
ACAGGCATAAGCCACCATGCCCAGCCTGCATTTATCCTTACATGATGGTGAAAAATAATG
TTTGTACTTCCTTCAGAATAATTTCAAGAAGGATCCCTGGAGTCAGCTAATGATTAGAGT
CAGGACTGTGCCTTAGTTGATGGCCCATATAGCACTACTGAACATGCCAGAGCTTTTGCT

16944    TTCAACTCTTACCAGCTCTATGAGCTTGAGCAGGTTACATACTCTTTTCAAGTCTTAGTG
CTTCACTTGTATTTTGGGGCTAATAAGGATTATACGAAATAATGCAGGTTAAATGCCTAG
CACTTTGCTTTACATACTAAGGGTTCCCAAGTGCTTTATTATTAGGTTTCTGAATGTTAT
ATATAAAGTTTCAGTGCTGCAAAAGGAATAGCACTCGAATATAACATTTTCTTTTTAATT
CTCAGCAAGGCAACGTACTTCTATATAGAAGGGTGCACCCTTACAGATAGAATAATGGTG
[G,A]
GCGCACACTTGGACAAGGGAGGAGAAGGGGTTCTTATCCCCCACGCACGTGGCCCCTGCT
CCTGTGTCGTTCCCCTATTGGCTAGGGTTAGACCACACAGGCTAACCTAATTCTGATTGG
CTAATTTAAAGAGAATGACGGGGTGAGGGCTTTGGCAGAGTCAGGGCAGAGCAGATAGCA
GGTAATCGGACTGAGTTAGGGTGGAGCAGGTGATCTGAATGAGTCAGGGTGGAGCAATCA
AAAAGGTTGCTTTATGAGGAAGTTACGTTTAAAAGTAGAAGGCAGGCTGGGCGCGGTGGC

17061    TAGCACTTTGCTTTACATACTAAGGGTTCCCAAGTGCTTTATTATTAGGTTTCTGAATGT
TATATATAAAGTTTCAGTGCTGCAAAAGGAATAGCACTCGAATATAACATTTTCTTTTTA
ATTCTCAGCAAGGCAACGTACTTCTATATAGAAGGGTGCACCCTTACAGATAGAATAATG
GTGGGCGCACACTTGGACAAGGGAGGAGAAGGGGTTCTTATCCCCCACGCACGTGGCCCC
TGCTCCTGTGTCGTTCCCCTATTGGCTAGGGTTAGACCACACAGGCTAACCTAATTCTGA
[C,T]
TGGCTAATTTAAAGAGAATGACGGGGTGAGGGCTTTGGCAGAGTCAGGGCAGAGCAGATA
GCAGGTAATCGGACTGAGTTAGGGTGGAGCAGGTGATCTGAATGAGTCAGGGTGGAGCAA
TCAAAAAGGTTGCTTTATGAGGAAGTTACGTTTAAAAGTAGAAGGCAGGCTGGGCGCGGT
GGCTCACGCCTGTAATCCCAGCACTTTGGGAGGCAGAGGTGGGCGGATCACGAGGTCAGG
AGATGCAGACCATCCTGGCTAACACGGTGAAACCCCGTCTCTACTAAAAATACAAAAAAA

17494    CTTTATGAGGAAGTTACGTTTAAAAGTAGAAGGCAGGCTGGGCGCGGTGGCTCACGCCTG
TAATCCCAGCACTTTGGGAGGCAGAGGTGGGCGGATCACGAGGTCAGGAGATGCAGACCA
TCCTGGCTAACACGGTGAAACCCCGTCTCTACTAAAAATACAAAAAAATTAGCTGGGCGT
GGTGGCAGGCACCTGTAGTCCCAGCTACTCAGGAGGCTGAGGCGGGAGAATGGCATGAAC
CCAGGAGGCGGAGCTTGCAGTGAGGCGAGATCCTGCCATTGCACGCCAGCCTGGGCGACA
[G,C]
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ATTAAGTCTTTGAAAAGAAATTTAGAACTCATATCTAACAATCCCTCCCCTTGTATTTCC
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17642    CTACTAAAAATACAAAAAAATTAGCTGGGCGTGGTGGCAGGCACCTGTAGTCCCAGCTAC
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[T,A]
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17737    GGAGGCGGAGCTTGCAGTGAGGCGAGATCCTGCCATTGCACGCCAGCCTGGGCGACAGAG
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[A,C]

# FIGURE 3JJJ

CATTTTTATGAGCCTCTGCATCTACTTACGGATGCACAGTATGACACAGCACCCGACAAG
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18068    GATGCACAGTATGACACAGCACCCGACAAGAATAAGTCCACCTATTACGGCTGCGAGGGA
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[A,G]
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18339    TCTGCTACTATCATGTCTAAGGCTATTTTTATTTTGCCAAGCCATCTGGCTAGTAGCCCCT
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[C,T]
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18361    CTATTTTATTTTGCCAAGCCATCTGGCTAGTAGCCCCTAATTGCTCAGCTATTCCATTAA
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[C,T]
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19218    CTTGTTGCCCAGGCTGGAGTGCAATGGCGCAATCTTGGCTCACTGCAACCTCTGCTTCCC
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[A,G,T]
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19298    TCTCAGCCTCCCGAGTAGCTGGGATTACAGGCATGCACCACCATGCCTGGCTAAGTTTGT
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[C,A]
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# FIGURE 3KKK

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19629    ATCTAATAGGGTTTGTCCCTGAAACCTCAGCCCCTATAGCATAAAACTGACTTAAAGAAG
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[C,T]
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19679    CTTAAAGAAGGGAACTGGCTTAGAAAAGGGGAAGAAATTTGAGAGTTTGAGATAATAACC
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19981    ATTAAATGGTCCCAATTCACTAATGCCCAGTCTGATGACAGTCAGGAGGCACAGAGGTAT
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[A,G]
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20014    GATGACAGTCAGGAGGCACAGAGGTATTTTTTTCTGAAATAGAGAGGTGTCTTTGACTTGG
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[C,T]
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20280    CGGCGCTTTCTTGACTCGGGTGTGATGAGTCCATCCCTTTTTCACTGTAGAAACAGCAGT
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CCACCCTTTGATAAGAACGTGATCTTCAGGCTGGTGTTGGTTTACCGGAAATTCTAGGGG
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[C,T]
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# FIGURE 3LLL

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20612 TCCTTTAGCACCTATTTTTATTAGATTTTTAGACCAAAGAAGGCCAAACACCATTTTTATAT
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[A,C]
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21966 GGTTTCCTCTAAAAGTTACTTTTCTACTTCCTTCTGTTAGCAAAGCAGTTGCGGCTACAG
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[C,T]
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22017 CGGCTACAGATTGAATGTATTCAGGCCATCCGCGGGTTACTGGGTTAAGGATTTTTGATA
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[T,C]
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28009 AATTTGTCCTTTTTTCACCCCTTCCTTGGCAAATCACGCAATATTCCTTCTTAAAAATGGGT
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[G,A]
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[T,A]
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# FIGURE 3MMM

28580 CCCAAATCCATCCTTTTTCTGTAGAACTTTAATGATGATACCTCATTCCTTTGTAACTTA
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[T,C]
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[A,C]
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28823 TGGCTGGGCGTGATGGCTCATACCTGTAATCCCAGCACTTTGGGAGGCTGAGGTGGATGG
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[-,A,C]
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28827 TGGGCGTGATGGCTCATACCTGTAATCCCAGCACTTTGGGAGGCTGAGGTGGATGGATCA
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[C,G]
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28842 ATACCTGTAATCCCAGCACTTTGGGAGGCTGAGGTGGATGGATCACCTGAGGTAGGGAGT
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[G,T]
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30128 GGCCAATGAATCTGAATTTAAAAACATGTATTTGTGTGATTTTGATGGGTGGACACACTT
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# FIGURE 3NNN

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30150    AACATGTATTTGTGTGATTTTGATGGGTGGACACACTTGAGAATCACGTCAGGACCATTT
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[T,G]
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30188    GAGAATCACGTCAGGACCATTTATGTGGCTCTCAATTACATATACACTACTTTATATTGC
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[C,T]
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30453    TTTCCTGCTACTTTCTCCCATTACAGGTCTCCCTACGGCAGCACTGCTTATATCACTTGG
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[T,C]
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34990    GGTTACACAGCTGGTGTCCTCGTTGCCTGTTCAGTAGAAAGGTTTACATAAACAGCAAGG
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AATTTTTAGAAAAAATAAAATCTTTTGCAAAGATTTTTACCTCAAAAATAGAAAAAAAGG
GCATTCCTGCCTTACCTTCTACAAGGGTCTTCTCTGAAATTCCAAGCATCAGGGTGTTAT
AACAGACTCTAAAAAGGGTTTCCTTTTTTTCTTTCCTTTAACATTGCTTATTGCACAGCAT
[A,G]
TTGAGACAGAGAAGATGGTAAGTGAAATAAAACAAAGGAAATAAAAAGTATCATCACTGG
GTTTCAGAATCAGCATGGTTTATGCTAAGGGAAAGACTTGGAAACCTTGATTCAACATAT
AATTCTAAAAAGAGACAGGAAGAAATCCCACCTTGTTTCCTCTGATTCTACCTTTGGGAT
GGGTAGGTATGTTATACAATAAGAATAACATTGAGATGACTGCTATAAAAATAGTGGTTA
AGAGCCTGGGTCCAGAATGAGAAAGGTGGATATTGAATTTACCTGAGTGCAACTAGGCAG

35203    CTGAAATTCCAAGCATCAGGGTGTTATAACAGACTCTAAAAAGGGTTTCCTTTTTTTCTTT
CCTTTAACATTGCTTATTGCACAGCATATTGAGACAGAGAAGATGGTAAGTGAAATAAAA
CAAAGGAAATAAAAAGTATCATCACTGGGTTTCAGAATCAGCATGGTTTATGCTAAGGGA
AAGACTTGGAAACCTTGATTCAACATATAATTCTAAAAAGAGACAGGAAGAAATCCCACC

# FIGURE 3000

TTGTTTCCTCTGATTCTACCTTTGGGATGGGTAGGTATGTTATACAATAAGAATAACATT
[G,A]
AGATGACTGCTATAAAAATAGTGGTTAAGAGCCTGGGTCCAGAATGAGAAAGGTGGATAT
TGAATTTACCTGAGTGCAACTAGGCAGACTCAAGTGAGTTGATTTTACCCACTCCTCCAC
TCAAATACTGGGTATGGCTTTGCAAAAACATTCAACCAGTTATCCACATAGTTGGTCTTA
ACTTTCCATGTGACTATAATGAATATAAACTTGCTAATGAGCAGAGTGTGATTTTAGTGT
TTAAACTATTTTTTCCCGAATAATAGTTCCTAGATGCAGTTAATGAGCCTTATTGGGTAC

36206   TCCAAGTAATTATCACAACATTAAGGTGCATTCAGCTTTGTGTGTTAACGTGGTATACCT
CCAGGCAACTTTTAGGATACTGTACAGATACAATGGCTGTGAAGGCTGGGATGAAAAGAC
CTGTGCGAAGCAGGACTGAGGCACTTAAGGAAGGCCTCAGAGTTACATCTCCTTTGCCTG
TTTTCTTGCAGGCCACATACCCTAGCCCAGCCCTGTCAGCATGAGTGAGAACCAGGCTCT
GCCTTTGCCCACACTAAACCACTACCTTCAAGGCCCCACAAAGACCCAGTGTCTCCAGAC
[G,A]
GTCTTTCTGTCTTCTTAACACTCAGAGCTCCATGAACCAGAATGAAAGTTTTGGAACATG
ATCCAAGTAAAAGACTCAAGAAGTAAACACCACTAAGGTTAACTTTGCTTTAGAGGTTAG
AGAAAACACTGCAAGGACACCACACCAGAGACTATGAAAACCCCAAATGTATTGAAATGA
TGCTGATTCCATTTACCTCCATATTGCCTGATAATACCCAGGTGCTACCATGGCAGCTTA
AGGTGGTATTTGCTGGGAGCTATGATACTCTTTAAGAAGTAATAGCACTACTAGTAAAAG

39692   AAAATTGAAATTACTTAGATACAAAAGAGTGGTTGTAGTAAGAAAATAGGCAAGGAGAAC
ATTTTAAAGTGCTGATCCTCGGTAAAGCCATACATAGGATGCACCTGGGAGCAGATCTTT
CTGAAGTCATTCTGTGCTCAGAGATGTTTCTCCTTACCTTGCTGCCTATGTCAAATTCTC
TGTGATATGTTCTTAGAGCCCCATGACCTCTCTTCTTAACTTGCAGTGGGAGCTTGAATT
TTCCATTTATTTTTGTGACCATTTAGTCTATAAGAGTCTCCGTCTTTACAGGGCCCTCAC
[C,T]
TGACTACAGACTCCATAAAGGCAGAGATTCTATTTTTACTCTATTATTACTGTATTCCCA
GCACTAAGCACTAGGATTAATACATAGTAAGTGTTCAACAGATGTTTACTGGATGATTAG
ATTGGCATTTTAAGGTAGTCTGAGATCACGTTTTAGACAAGATACTTCAGTTTAGTCCAA
TCTTTATTATTTATTAGCTACTAAAGAGAAATTGATAATTACTCATGATATTCTTCTTTT
TTGTTTTACAGTCAACTTTGACCACTTTGAAATTTTGCGAGCCATTGGGAAAGGCAGTTT

40095   TGTTTACTGGATGATTAGATTGGCATTTTAAGGTAGTCTGAGATCACGTTTTAGACAAGA
TACTTCAGTTTAGTCCAATCTTTATTATTTATTAGCTACTAAAGAGAAATTGATAATTAC
TCATGATATTCTTCTTTTTTGTTTTACAGTCAACTTTGACCACTTTGAAATTTTGCGAGC
CATTGGGAAAGGCAGTTTTGGGAAGGTGAGAACAAATTGAAATGATTAACCACCAGCAGG
GTTATGTAGCCCAGGGAACAGAGGGTCCAGAAATGTTCACATTATTGAGTTGCTGGGACC
[A,G]
CAAGGAAAGATAATTAAGTGAAAATGTTTTTGTAATGGATTTTTATAAAATTGTCACCAC
AGTTTAAGAAAAGCGTGTGACAGGCAGCTACATAATGAACATATACTGTTGTCAGAATAA
TCTCATTAAACTCAAATCTGTTTACTCTCAGTAAACTTTAAGGCTTTTCTCTCTACCCTA
AAGGAGATGAAGATTTCAGAATCATTTTCAGATTCTACCAGCTGTATGCCCAGTAATAGT
TATCTTGTTTATGGAAGAGTTACTTATTTTCATGTGGGAAAGAAGTCATCCGATTTCTAT

40191   TACTAAAGAGAAATTGATAATTACTCATGATATTCTTCTTTTTTGTTTTACAGTCAACTT
TGACCACTTTGAAATTTTGCGAGCCATTGGGAAAGGCAGTTTTGGGAAGGTGAGAACAAA
TTGAAATGATTAACCACCAGCAGGGTTATGTAGCCCAGGGAACAGAGGGTCCAGAAATGT
TCACATTATTGAGTTGCTGGGACCACAAGGAAAGATAATTAAGTGAAAATGTTTTTGTAA
TGGATTTTTATAAAATTGTCACCACAGTTTAAGAAAAGCGTGTGACAGGCAGCTACATAA
[T,C]
GAACATATACTGTTGTCAGAATAATCTCATTAAACTCAAATCTGTTTACTCTCAGTAAAC
TTTAAGGCTTTTCTCTCTACCCTAAAGGAGATGAAGATTTCAGAATCATTTTCAGATTCT
ACCAGCTGTATGCCCAGTAATAGTTATCTTGTTTATGGAAGAGTTACTTATTTTCATGTG
GGAAAGAAGTCATCCGATTTCTATTTGTTTCCTCATTTGTCTAATGTTTTTATCTTAAGA
AAAATACATATTCAGTTTAATTTTTTTTGCAAGAAACTTCTGTATTCAAACCCTGATTAC

40287   CAGTTTTGGGAAGGTGAGAACAAATTGAAATGATTAACCACCAGCAGGGTTATGTAGCCC
AGGGAACAGAGGGTCCAGAAATGTTCACATTATTGAGTTGCTGGGACCACAAGGAAAGAT
AATTAAGTGAAAATGTTTTTGTAATGGATTTTTATAAAATTGTCACCACAGTTTAAGAAA
AGCGTGTGACAGGCAGCTACATAATGAACATATACTGTTGTCAGAATAATCTCATTAAAC
TCAAATCTGTTTACTCTCAGTAAACTTTAAGGCTTTTCTCTCTACCCTAAAGGAGATGAA
[G,A]

# FIGURE 3PPP

ATTTCAGAATCATTTTCAGATTCTACCAGCTGTATGCCCAGTAATAGTTATCTTGTTTAT
GGAAGAGTTACTTATTTTCATGTGGGAAAGAAGTCATCCGATTTCTATTTGTTTCCTCAT
TTGTCTAATGTTTTTATCTTAAGAAAAATACATATTCAGTTTAATTTTTTTTGCAAGAAA
CTTCTGTATTCAAACCCTGATTACTAGTTTCTCAATGGAGACGTACTTTAAGAGAATAAT
ATTTCATATAAAACTTGCATTTTAAAATCATTTTCTGTTTACTTTTTCAGGCATTATACA

40384    TTGCTGGGACCACAAGGAAAGATAATTAAGTGAAAATGTTTTTGTAATGGATTTTTATAA
AATTGTCACCACAGTTTAAGAAAAGCGTGTGACAGGCAGCTACATAATGAACATATACTG
TTGTCAGAATAATCTCATTAAACTCAAATCTGTTTACTCTCAGTAAACTTTAAGGCTTTT
CTCTCTACCCTAAAGGAGATGAAGATTTCAGAATCATTTTCAGATTCTACCAGCTGTATG
CCCAGTAATAGTTATCTTGTTTATGGAAGAGTTACTTATTTTCATGTGGGAAAGAAGTCA
[T,C]
CCGATTTCTATTTGTTTCCTCATTTGTCTAATGTTTTTATCTTAAGAAAAATACATATTC
AGTTTAATTTTTTTTGCAAGAAACTTCTGTATTCAAACCCTGATTACTAGTTTCTCAATG
GAGACGTACTTTAAGAGAATAATATTTCATATAAAACTTGCATTTTAAAATCATTTTCTG
TTTACTTTTTCAGGCATTATACAGACCTCTAAAGAAATTTCAAAAACATGGACATCATAT
TTAGTGTTTTTCCAGTCCTTAAAGTCCTTTTTGGTTATATCATGTATGGGTTGTAAACAG

40510    GAATAATCTCATTAAACTCAAATCTGTTTACTCTCAGTAAACTTTAAGGCTTTTCTCTCT
ACCCTAAAGGAGATGAAGATTTCAGAATCATTTTCAGATTCTACCAGCTGTATGCCCAGT
AATAGTTATCTTGTTTATGGAAGAGTTACTTATTTTCATGTGGGAAAGAAGTCATCCGAT
TTCTATTTGTTTCCTCATTTGTCTAATGTTTTTATCTTAAGAAAAATACATATTCAGTTT
AATTTTTTTTGCAAGAAACTTCTGTATTCAAACCCTGATTACTAGTTTCTCAATGGAGAC
[G,A]
TACTTTAAGAGAATAATATTTCATATAAAACTTGCATTTTAAAATCATTTTCTGTTTACT
TTTTCAGGCATTATACAGACCTCTAAAGAAATTTCAAAAACATGGACATCATATTTAGTG
TTTTTCCAGTCCTTAAAGTCCTTTTTGGTTATATCATGTATGGGTTGTAAACAGAAATTC
TTTGCACAGTATTATTCAGCTTGACAGTTCAGTCATGTCTATTTCAGTCACTCAAAGCAG
GATTAAGGATGTTACTTGTTATTGGAATATTCCTGACATGGAGGCAGCTATTTTCACCAA

41664    GAAAACAGTGAAAGTACAGAGATTTATTGCAAAGTGGAAAAGTACACACTCAAGAGAGGG
GAGCATGGGTGAACTCCAGCGAATGTCATGTAAGGGGGGGTTTGAGGCTGCTGCCATAAT
GGGTTTCTTTAACCAAGGGGTGAAACATTCATGATGATTCCTGAAAAAAGATGGAGATTT
CTTGGAACTGTGGTGCCAGCTATTTTTACACCAAATATGAATGTTCCTGGAACTGTCATG
GTGCTGGTGGGTGTATGATTTAGTATGTTAATGAGTGTATGATGAGGTCCTAGGTGAAAC
[C,T]
TAGGTCAAATCCAGCACAATGGAGAGGACCCACAGACTCTCTGAAGGAAACGACTGCTCC
TGCAGGACCCAGGCAACTCCCCCAAAACTGTGAGTACCCCAACTGTGGAGGTGGGAAAGA
GAGACCCTCCTCTCCCAAACACACACCCCCACTGGAGAAGCTGAAGGTCTGTTTGCTGGA
GAAGTTTCTGACTTTACCTGGAGCTGAGTGGACTTGAAGAGCCCAGTGAAATACACGGGG
AGAAGAAGCAGCAGAAAGGCCCTGGGAGCTTGCTGGGTCCACAAGCAGGCCATTCCTGCC

48324    CACTGATATGTGGAAGCTAAGCTATGAGGATGCAAAGCAATGAGAATGATACAATGGACT
TTGGAGACTTAGGGGGAAGAGTGGGAGGGGGGCGAGGGATACAAGACTACAAATGTGGTG
TAGTGTATACTGCTCAGGTGATGGGTGCAACAAAATCTCACAATCACCACTAAAGAACTT
ACCCATGTAACCAAAACCACCTTTACCCCAATAACTTATGGAAAAATAATCCAGCACCAC
ATTAGGTTTAGTCGGACTTAGCCAGCTTGGCTTACACCCTGGTTTTTCAGGTTCTTATCA
[T,G]
TCCCAGTTTATGCAGCTGTTTCAACATTTTCCTTTTGCTAGTCATGTGAAACTGCTGTCT
GGAATTTTCTTTTCTCCTGCTACCACCCTTTATTATTCCTGTCTCACTTTCATCTTCATC
CCTACTGTTACATAAATGCATCTTGATTTCTAGGCAAGCATTTGTCAAATTCTCATTAGG
ATCTTCCTCAGGGTCTTTTGTTCTCCTTAGTTTCTTTGGCTTTATAGTGAAAGAACATTT
TTCTTTTATTGTCACTAACAAATACTTCTTGGTCAGTTGTCACAGTTCCCCTTGTCCTTG

48423    TACAAGACTACAAATGTGGTGTAGTGTATACTGCTCAGGTGATGGGTGCAACAAAATCTC
ACAATCACCACTAAAGAACTTACCCATGTAACCAAAACCACCTTTACCCCAATAACTTAT
GGAAAAATAATCCAGCACCACATTAGGTTTAGTCGGACTTAGCCAGCTTGGCTTACACCC
TGGTTTTTCAGGTTCTTATCATTCCCAGTTTATGCAGCTGTTTCAACATTTTCCTTTTGC
TAGTCATGTGAAACTGCTGTCTGGAATTTTCTTTTCTCCTGCTACCACCCTTTATTATTC
[C,T]
TGTCTCACTTTCATCTTCATCCCTACTGTTACATAAATGCATCTTGATTTCTAGGCAAGC
ATTTGTCAAATTCTCATTAGGATCTTCCTCAGGGTCTTTTGTTCTCCTTAGTTTCTTTGG

# FIGURE 3QQQ

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        CTTTATAGTGAAAGAACATTTTTCTTTTATTGTCACTAACAAATACTTCTTGGTCAGTTG
        TCACAGTTCCCCTTGTCCTTGAGGTCAATATATATATATTTTTAAACATTGTAATTAAAT
        ATGCTGACTGGGAAGGAGTTCAGATGTCTTACTAGTTATTAGATACTTTCTTTCCCCATG


50015   GCCTTAGCATTTTACCTTCTCATATTTGTCTTTCATCGCTGTGTGGGCAAAGTTGATTTC
        ATTCTGTTCCTTTTTTTTAAGAAAATGGGTATTGTGAGGCTTTAAGCTGGCCAAAGATGAT
        AGATTTTGCTGTTTGCTAATTTGGTGTCATTCCAGACAACATTCTGTTCTCCATGCATAC
        TGACCTGGTGATAACATGACATATAACCTATTCTTTCCTTCTCACTTCTCACATTGAACC
        TCACAGTGGAACACTAGGCATCATTAACAATGATAGAAGAAAGAGAGGAGACTTACCTCC
        [A,C]
        CCCAGTGATTCTGGTACTACATTCAAAACTAGAAACTAACTGGGAGGGGGAATTCTTAAA
        GTACAACAGCAACTCCCTTTGTCTTCCAAACCATGAGAAAAATCTTCACAAATCTGTATC
        ATTCTTCCTAATAAATGCTTTTTGTTTTAGTAAGTACAATATATTCAATGTAAGTTTATC
        TTTCCACATTTATAAACCATCTTGCAGTGCTTTTGAAGGTGTGATTGTGAGTGTATTAGT
        CAGTTCTCACATTGCTATAAAGAAATACCTGAGACTGGGTAATTTTTAAAGAAAAGAAGT


50095   AAAATGGGTATTGTGAGGCTTTAAGCTGGCCAAAGATGATAGATTTTGCTGTTTGCTAAT
        TTGGTGTCATTCCAGACAACATTCTGTTCTCCATGCATACTGACCTGGTGATAACATGAC
        ATATAACCTATTCTTTCCTTCTCACTTCTCACATTGAACCTCACAGTGGAACACTAGGCA
        TCATTAACAATGATAGAAGAAAGAGAGGAGACTTACCTCCACCCAGTGATTCTGGTACTA
        CATTCAAAACTAGAAACTAACTGGGAGGGGGAATTCTTAAAGTACAACAGCAACTCCCTT
        [T,G]
        GTCTTCCAAACCATGAGAAAAATCTTCACAAATCTGTATCATTCTTCCTAATAAATGCTT
        TTTGTTTTAGTAAGTACAATATATTCAATGTAAGTTTATCTTTCCACATTTATAAACCAT
        CTTGCAGTGCTTTTGAAGGTGTGATTGTGAGTGTATTAGTCAGTTCTCACATTGCTATAA
        AGAAATACCTGAGACTGGGTAATTTTTAAAGAAAAGAAGTTTAAGTGGCTCATGGTTCTG
        CAGGCTGTGCAGGAAGCATAGTGGCTTCTGCTTTGGGGAGGACTCAGGAAGCTTCCAATC


52300   GTATGAAGAACTGTGTGTGTGGTGTGTGGGTATGTATGTGTTGGGGTTTCAGAGAAAGAA
        GGTAAGTAGTCTGGGGGCAGGGACGTTAAGGAGGAAAGAACATTTGGAAATAAAATTCAA
        CCTGACTTGCCTCCAGGGACCTGGCTACACTCAGGAACAGTCTTCAAATGTAGGCCATGT
        TATCAAGTGAATGCTGCCAGACAGGGCTGGCATCCAGGAAAAGTAAATAAAATCTTCTTG
        TGCGTCTGTCTCTGAGGGCTCTTCACAAAGCCCTGGCAACCCACAGCCTGAAAACAAATA
        [A,G]
        GCCCCAGTCTTTCCCAGCATAGTTGATTCCCCAGGTGGCTTTTGTTAATTGAGATTAAAC
        CTGTAGCTGCACACAACTCCTCAGGGCCTCTATCTCTTTACTCATGTCTTTGTCCCTGTG
        GATAGAAGGGGTCCACATGTGGTTTCAGGAAATTAGGACACCAGATCATCTGTTTTAACT
        GGAAAGAACTACCTGTACTGAGAGTGTGACAAGGTCCTTTCAGACTCTGAACATAGCCCA
        ATAAATGGTATCAACCTTAAATAACGAGATTCTGAAAATATGATTAAGTATCGAGTTTGC


52623   TTGATTCCCCAGGTGGCTTTTGTTAATTGAGATTAAACCTGTAGCTGCACACAACTCCTC
        AGGGCCTCTATCTCTTTACTCATGTCTTTGTCCCTGTGGATAGAAGGGGTCCACATGTGG
        TTTCAGGAAATTAGGACACCAGATCATCTGTTTTAACTGGAAAGAACTACCTGTACTGAG
        AGTGTGACAAGGTCCTTTCAGACTCTGAACATAGCCCAATAAATGGTATCAACCTTAAAT
        AACGAGATTCTGAAAATATGATTAAGTATCGAGTTTGCTGGAGCCCAGAGCTTGAGGATG
        [C,G]
        CCACCTGGGAGCACAGATTCACTTTGCCCAGAATGTACACTCCAATTAGCAGCAGTTATA
        AGTGGGGTTTTAAGAAAAAAAGACAAGGCAGTTCCTAAGTTATTTACCAAAAATTTACAT
        TAAAATAATGTAAGCTATTGATGGACTATGCATTATTCTTTATATCACAAATTACAGGAA
        CACAAAGATAATGGGTGAGGCAGCTAGTCAGGAACAAAATGGCTTTAAAAATACTGTCCTT
        GAGCATGGGTTTGAGGCTGTGACTGACATCCCATACTCATGTTTCTCTAAACCTAATAAA


52773   TTTTAACTGGAAAGAACTACCTGTACTGAGAGTGTGACAAGGTCCTTTCAGACTCTGAAC
        ATAGCCCAATAAATGGTATCAACCTTAAATAACGAGATTCTGAAAATATGATTAAGTATC
        GAGTTTGCTGGAGCCCAGAGCTTGAGGATGCCCACCTGGGAGCACAGATTCACTTTGCCC
        AGAATGTACACTCCAATTAGCAGCAGTTATAAGTGGGGTTTTAAGAAAAAAAGACAAGGC
        AGTTCCTAAGTTATTTACCAAAAATTTACATTAAAATAATGTAAGCTATTGATGGACTAT
        [G,A]
        CATTATTCTTTATATCACAAATTACAGGAACACAAAGATAATGGGTGAGGCAGCTAGTCA
        GGAACAAAATGGCTTTAAAAATACTGTCCTTGAGCATGGGTTTGAGGCTGTGACTGACATC
        CCATACTCATGTTTCTCTAAACCTAATAAATTGTGCATATCTCATATAGCTCAGACTGCT
        CTGAGCTATTTTTGTTTTCTCATTTCCCCCCTTTTCATCAAGATTTTGCAAAGAAAGCAT
```

FIGURE 3RRR

TGTGGATGAACTTAAGCAGTTTTGGCTCCTTTTATGTTCAGGAACTTAGTCCTGCATTGC

53140   AAATGGCTTTAAAATACTGTCCTTGAGCATGGGTTTGAGGCTGTGACTGACATCCCATAC
TCATGTTTCTCTAAACCTAATAAAATTGTGCATATCTCATATAGCTCAGACTGCTCTGAGC
TATTTTTGTTTTCTCATTTCCCCCCTTTTCATCAAGATTTTGCAAAGAAAGCATTGTGGA
TGAACTTAAGCAGTTTTGGCTCCTTTTATGTTCAGGAACTTAGTCCTGCATTGCTAGGAA
GTCTTATTCCCAGATGGTCCTGTCCCACATTTGGGGGAAGGGGAAAGGATGAGTCTTAGT
[G,A]
GGGATTTTAACACCATCAGAAGCAAAATTGGGATGGCATCGCAGGGTGCCACAAATGAGA
CCTCACCCAAGTCACTAATTTATGTAGCTACTGTTGCTTGTGGGATCATCTCCAGGCTTC
AGAATACCATGCAGTTAGTTTTCTCGGAATAAGTAAAACAATGAGCTATACATAGTAGAA
ATATAATACACATAACAATTACAATTAAAAAAAAAAAAAGAATTTCTATGCCTGAATGAAA
AAAATATCTATTCCATTGGAAAGTCAACTAAAAACATCATGAAGAAAATTAAAATCCAGT

53848   GTGTGCTATAATTTTCTTCTGAACCATAATTTCTCTCTCTTCAGTTCACTATTTCTACCC
AAGATAAATGTTATCAGGACCAACATACTTGTAAAATAAGCTTTAGTATTATATTTGGCC
TAATTATTTGCATTAAGTGCAACAAAAATAATGAATGGCCATGTACGCATTTTTAAGTTG
GCTTTGCTGGAACTTTTTCATAAGGAATCTCAGATTAGACTTTTAAAAGCCTCTCTAAAC
TAGATATTGAAGCCAATAATTCACCATCAAACTGCCTGTAGCATCTACATAAATTGGGTG
[A,T]
ATTTCTCCCTTCTTCAGGTTCTGAAATATATTGAGGTTTCTAGGCCTGTCAAATGATGAC
ATTCTTTACTTACTGCAAGGTCAAAAAACTTGTGAGGGTACCATGTAGACAAGGTATCAG
GTCAGTTTTCCAAAAGGACTATTGATTTGGCTCTATAAAGTCAACTTCAATTCATCAAAG
CAGTTTGGTCATATCTGAAAGTATGTCATTTCACCCAAAGCCTTGGTAAAATGACCAGCC
TTAGTAAAATGACCAGTGTCTCCAACTGTGTACTGTTACAGAAGAAAACAGGTTCTTACT

57636   TATGTGAACTAAAAGGGATTTGAGTTATTTTCTATTTTTCTGATAAAATATTTAAGTGTT
TCCTTTCTCTTTTGGCCAATTAGAACTCATTCATATATTTTTGTAATAAATTTTACATAC
ACATGACACATATAAACATGCAGACACACACAGGCAGATTTTATAGCTTTGTAAGTTTCT
TCATTTGCCAGTTTTCAATAGTTTCTCTCCCACCTTTAGACTGTCAAGCCCTAAACAATT
GTTAGCTAGGCAACCTTAAATTTGTACTTCTAAAGGGATGACTCTTAGCTGAAACAAAGT
[-,A]
AAAAAAAAATAAAAATTACACTTCAAAAACACAGAGCGGAGCTCAAACTAAGGGAGCAGGT
GTATATAGGTAAAGGTCCAGTTAAGACAAGATGGCCAAGGAAAGCATCTTAAGTAAAGGT
AGGACTTGTATAGATTTAAACCAATGTTAAAATTTCTCATGACTCAGCTCTCCCTCTCCTC
CAGGTGCACAGAGGCAGAAACCCTTACAAATGGAGATTTCCTTTATCAATGTAAATTTCA
ATATAGCCAGCTAAATGCCAGCAAGGTATATTTTGGAGAACTGTTAGAGGCAGTGAATCT

57693   GTTTCCTTTCTCTTTTGGCCAATTAGAACTCATTCATATATTTTTGTAATAAATTTTACA
TACACATGACACATATAAACATGCAGACACACACAGGCAGATTTTATAGCTTTGTAAGTT
TCTTCATTTGCCAGTTTTCAATAGTTTCTCTCCCACCTTTAGACTGTCAAGCCCTAAACA
ATTGTTAGCTAGGCAACCTTAAATTTGTACTTCTAAAGGGATGACTCTTAGCTGAAACAA
AGTAAAAAAAAAATAAAAATTACACTTCAAAAACACAGAGCGGAGCTCAAACTAAGGGAGC
[A,T]
GGTGTATATAGGTAAAGGTCCAGTTAAGACAAGATGGCCAAGGAAAGCATCTTAAGTAAA
GGTAGGACTTGTATAGATTTAAACCAATGTTAAAATTTCTCATGACTCAGCTCTCCCTCTC
CTCCAGGTGCACAGAGGCAGAAACCCTTACAAATGGAGATTTCCTTTATCAATGTAAATT
TCAATATAGCCAGCTAAATGCCAGCAAGGTATATTTTGGAGAACTGTTAGAGGCAGTGAA
TCTGTATGTGTCTGCAGCAACTTCAATTCTTGCCTACTCTCAAAATAAAAAAATTCAACTG

58585   GAAGTTATGTGCATGCTTACTTGAGGCATCTTTTTTTTCCTTACCAGTTGACTGTTCCTAG
AGGAAGGTCATATACCAGTTAAACTCTACCATTTTTGCCTCTTAGTGTGCATGCTTGAGC
CTACTCGCCCACCTCCTGAGATCTTATCAGGAACCTACTGATCATCAGTTTCAGGGTTTT
TCTATCTACTGGGAGATTGCCTTTTCCTGGCGCCGGCTGCAACCAAATATTATTTGAGAG
AGACAGTTTAACAACCACCTGACCATCACCTAATGGTTGTCTGACATTCCTTGGTGGAGG
[T,C]
TGGGGGTGATCTCCTGCCTTGCCCATGTCTGCCTGCCTACTGTAACAGACCAACTTAGTT
AAATAGGTGGGCTTTTTCAACTTAGTTTGTTTCTTGGTGAGATGACTGACATCATTGTGAA
GCTCTTTAATGAACAGGGCAAAGAAAGCCTTCTCTATGCCTGGACTCGGCATGGACAGCT
CTGGGAAAGAAGAAAGCCTATTTTACCTGAGGGCCTATCTTTTATAAATATTTTGTTCAA
ATTCTTTCTTTTAAAACAAAGGTTCTTTTTTCAATGACTTACCAAACCAATACACCTTAAC

FIGURE 3SSS

58649   AGGTCATATACCAGTTAAACTCTACCATTTTTGCCTCTTAGTGTGCATGCTTGAGCCTAC
TCGCCCACCTCCTGAGATCTTATCAGGAACCTACTGATCATCAGTTTCAGGGTTTTTCTA
TCTACTGGGAGATTGCCTTTTCCTGGCGCCGGCTGCAACCAAATATTATTTGAGAGAGAC
AGTTTAACAACCACCTGACCATCACCTAATGGTTGTCTGACATTCCTTGGTGGAGGTTGG
GGGTGATCTCCTGCCTTGCCCATGTCTGCCTGCCTACTGTAACAGACCAACTTAGTTAAA
[T,C]
AGGTGGGCTTTTCAACTTAGTTTGTTTCTTGGTGAGATGACTGACATCATTGTGAAGCTC
TTTAATGAACAGGGCAAAGAAAGCCTTCTCTATGCCTGGACTCGGCATGGACAGCTCTGG
GAAAGAAGAAAGCCTATTTTACCTGAGGGCCTATCTTTTATAAATATTTTGTTCAAATTC
TTTCTTTTAAAACAAAGGTTCTTTTTTCAATGACTTACCAAACCAATACACCTTAACCAAG
GTTATGTCTAAACCAAGGATCAACTAGGCATTTCCAAAGAGTGGCAAAGTAGTCCTCACA

62188   GGTGGCCTTCAGAGCCACAGCATCAACAATATTAACTTCCCTATTAGTAGTGTTCTATTA
CTTTGGGTTTTACATATATTATCTCATTTATTCATCATAACAACCTGGTTGATAGGGATT
ATTATTCCCATTCTATTCCTGAAGAAACTGAGGCTCAAAGGAGCTAAAATATTTTCCTAT
AGTCACACAGCTAGGAAGTGGCAGAGCGAGGACTCAAACCCAAGAATCCTGACTTCAAAG
CCTCTGCTCTTCCTGCTGCACTATACCATCCCTATACACATCTCTGAGACTCCTGTAAAA
[A,T]
TATGTAAGGAACAGGATTTATTTCATTTATTGTCTTTCATATCCCACAAGAATACAAACT
GTGTAAGGCAGGTATGTCTGTATGTTTTTTATCACTGCCTCATTCCCCATCTTCCACAAC
AGTGCCTACCGCACAGTAAGTGCTCGATAAATATCTTTTAAATGAGCATGTGAATGAATG
TGTGTTAGTGTTAGGGCTAAGGCCTTTGGCTTCTGGTTAATTGCCCTTTTTGCCATTATG
CCAATGTCATTTGCACACTCACAAACATACCCTCATATAATCATATGCACTTCAGTTTCT

63478   CATCTCAAGATATGTCAAATAAGTGTATTTGGGGTGAAATATTTTTGGTTTCCTTTGCTA
GAAATGAAATGTCCCTGCTTCCCCATAGCCAGAAAAGATTCTTGAGTGGACAACTGCACC
TAAACTTGAACCTGAGCACTAGAAAGTCTTTTGTTTTATTCTATGTTTTTATAAATTTAA
ATCTAATTTTTTGAATATAAAATAATACATATTTTGTAAATGTGGAAACACAGAAAGTTC
TAATGAAAAAATAAAAACCTGTATTTCATCACGCAGAAATATCTGCTGTATTAGTTTTCC
[G,A]
TTGCTGCGGTAACAAATTGCCACAAACCTGGTGGCTTGAGACATCATAGATTTAGTATCT
TACAATTCTGGAAGTCAGAAGTCCAAAATCAGTCTCCCTAGGCTAAAATCAATGTGTCAC
CAGGGCTGTGTTTCTTCCAGAGCCTCCAGGTGAGAATCTGTTTCATTATCTTTTCTAGCT
TCTTGAGGCTGCCTGTATTCTCGGCTTGTGGCCCCTTCCTTTATCTTCAAAGCCAGCAGC
ATACTATCTTCAAACCTCTCTCTGACTCTGACTTCATGTTCTCCTTATTCATCTTTTAAG

65457   AACCTTTTTCTTCTGCCATGAGACTAACCCTGGCTTCTTCACGTGCGGGTGGAAGGGTTC
CTAACAGCAACAGCTGACAAACTTAATGAGCAAGCACTTTTTCAGCCTCTGCCACAGTCA
CATTTTCTATCCTATTGGCTAAAGTAAATCACGAAGTCAGGCTCAGATTCAAGGGGTGTA
GAAATAGGCTCCACTTCTGATGAGTGGCACGGCAAAGTCAACATTGCAAAAAGCCAGGCA
GAGATATTACTGTGGCCAGTTTTGCAAACAATCCACCGTAATACATAAAATATGTTTAAG
[C,A]
AGTCCACAAAATGATCAAGGAAATGGTAGAAACTATAAACACTGCAAGAACTCAGAGCCA
CATGATGTTATTGAGTCCTTGTAGTGCTCTGAAAGGGTTCAAGGAAGAAGTTGTTTTGGC
ATATGACCCTGATGAACTTGCAAAAGTAGAGAAGAAGGGAGCACAGTTTCTGAAGAAGAA
CTTAGTAGAGAAGTGTTATTCTGTGGCCAGTACGCAGTAATTGTTCCACCTAGAGATGTT
GACTGACTGATGAACAGGAAGCTGAGTCTTTATAATGCAGATATTCACATATTCATTTAC

69947   CTGTAAATCCCAGCTACTTGGGAAGCTGAGGCATGAGAATCCCATGAATCCTAGAGGTGG
AGGTTGCAGTGTGCCGAGATCATGGCGCCAATGCACTCCAGGTTGGGCGACAGATCCAGA
CGCTGTCTCAAAAAAAAAAAAAAAAAAAAAAAAAATCTTTGCCTATGCCAACGTGGAGCTA
TTCTATCCTGTTTCCTAGAAGCTTCACTGTTTTAGCTTTCACATTTAGATCTACAGTCTA
GGATCAAGTTTTATTTTGTCTTCATATAAATAAGTAATTGACCCTTAGCCATTTGTTGAT
[A,G]
AGCTTATACTTTCCTTACGTCACCACAGAACCACATTTGTTATTAATCAAGTCACCATCT
ATGTATGGGTTTCCTGACTCTGTTCCATTGATTCATTTGTATACTCTTGCATATTTATCA
CTCTGTTTTAATTACTGTAGTTTTTATACTGGATTTTTCAGTAATTCATCTTTGGATTATGT
TGGCTACAGTTGGTTCTTTAAAAATTCCATATAAATTTCATAAGTAGCTTTTCAATTTGTA
TTTTAAAGCTGCTGGTATGTATATTGGGTACATGGAGTCTATAGATTAATTCAGGGATAA

69981   GAGAATCCCATGAATCCTAGAGGTGGAGGTTGCAGTGTGCCGAGATCATGGCGCCAATGC
ACTCCAGGTTGGGCGACAGATCCAGACGCTGTCTCAAAAAAAAAAAAAAAAAAAAAAAAAA

FIGURE 3TTT

```
      TCTTTGCCTATGCCAACGTGGAGCTATTCTATCCTGTTTCCTAGAAGCTTCACTGTTTTA
      GCTTTCACATTTAGATCTACAGTCTAGGATCAAGTTTTATTTTGTCTTCATATAAATAAG
      TAATTGACCCTTAGCCATTTGTTGATAAGCTTATACTTTCCTTACGTCACCACAGAACCA
      [C,T]
      ATTTGTTATTAATCAAGTCACCATCTATGTATGGGTTTCCTGACTCTGTTCCATTGATTC
      ATTTGTATACTCTTGCATATTTATCACTCTGTTTTAATTACTGTAGTTTTATACTGGATT
      TTCAGTAATTCATCTTTGGATTATGTTGGCTACAGTTGGTTCTTTAAAAATTCCATATAAA
      TTTCATAAGTAGCTTTTCAATTTGTATTTTAAAGCTGCTGGTATGTATATTGGGTACATG
      GAGTCTATAGATTAATTCAGGGATAACTAACATCTTTTTAAAAATATCAAATTTCCAATTC
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71165
```
      ACTAGAAGTTTTTTTTTTTTTTTTAACATGAATGAGTACAATATTTTATCAAATACTTTTGT
      TTTACTGAGGTCATTTCTATTGTGAGTGAAGCAAGTTGATTTGTAAATATTAAAGCAATC
      TTGATTTCCAAAAGTAAATGCTAGTTGGTCATGTTCTATTATCCTCTTGTGTATATTACT
      GGCTACAATAAAATATTTGTTTTTTTATATTTTTTATATTATTATTCATACATTATTTATG
      TATGTTATTTATTATTTATAAATATGTATTCTATTTATATATATTCCTACATATATTTTA
      [G,A]
      GATGTACATAGACAAGTTTGAATGGTAACAAGAATGAGCCAACTGAGAGGAAGAAATTGG
      TAATGTAGTAAAGAGCGGGGATGATTGCCAAGTCAGGTCCTGCAGGTGGTGAGATGAATG
      TGACTCAGGGCACAGGTGAATGAGCTGACCTTAGGTGGAAGTGGGGACCCTTCCTTCATG
      TACTAGGAGAGAAAGCAGAGTTTGAAGTCTGTATGTGTGTGAGCTGCTGGGCTTCTCAGA
      GGGCAGATGAAATAGTTCTTATGCCATTGCCTGTGTTTTCCCTGTGGTATATGAGGCCAT
```

71347
```
      CTACAATAAAATATTTGTTTTTTTATATTTTTTTATATTATTATTCATACATTATTTATGTA
      TGTTATTTATTATTTATAAATATGTATTCTATTTATATATATTCCTACATATATTTTAGG
      ATGTACATAGACAAGTTTGAATGGTAACAAGAATGAGCCAACTGAGAGGAAGAAATTGGT
      AATGTAGTAAAGAGCGGGGATGATTGCCAAGTCAGGTCCTGCAGGTGGTGAGATGAATGT
      GACTCAGGGCACAGGTGAATGAGCTGACCTTAGGTGGAAGTGGGGACCCTTCCTTCATGT
      [A,G]
      CTAGGAGAGAAAGCAGAGTTTGAAGTCTGTATGTGTGTGAGCTGCTGGGCTTCTCAGAGG
      GCAGATGAAATAGTTCTTATGCCATTGCCTGTGTTTTCCCTGTGGTATATGAGGCCATCC
      ACTGAGAATGAAGGTGGTCAGAGTATAGGAAATTTTGAGATGCCGAGAAGATCTGTGAAA
      TTAGTAGAGAATTAGAATAGGATTTTCTAAGTATCCATTTGAGACTTGTAGTTATAATTA
      AACAAGAATCTATCCTGCAGATTTGTATTTTTCTCCTTAGATTGCACTTAATAGATCACC
```

71903
```
      TGCAGATTTGTATTTTTCTCCTTAGATTGCACTTAATAGATCACCAGTTCATTTTTGTTG
      CTGTTTAAAAGCATATTGAGTTTAAGCAGGATTGGAGTTTAATTGGGTGAGGTATTCTCA
      CTGTGACTAAGTTTGATGAATTGAAAAGCGTAGTTGTAGAAAGGAAACTCAAGAAGGAAA
      TTCTTGGGGAAACTTAAAGAATCGTATATATGCAATGTCACTTTTTAAGACAACTAATAT
      TTTTAAGAATTTACTACTTTTGAGGTGCTGTACTAATATATTACATGTATAATTTCATAT
      [A,G]
      TCTTCAACTACTAGTTCCTGTAAATAAGTATGCTGATGATGACACGTTCCATTTCTTTCG
      ATAGCCACACAAAAACAGGAAGTGATGACAAAGCTGGATTCTAACTCCTGACTCCCAAATTC
      TCTAAGACCCTCAGCATTAACATATATTTTATTTTAATGTTATTATATATGTATCATTAC
      TTTTACAACTCTTAAACCAAACATTTTAAAATTAGCTACAACTGCAAAATCAACTTAAAA
      ATTTCAAAGAGCCATTTAACATGATAAATTAAAATATTTTAGTAAAACAAAATCACCACT
```

71908
```
      ATTTGTATTTTTCTCCTTAGATTGCACTTAATAGATCACCAGTTCATTTTTGTTGCTGTT
      TAAAAGCATATTGAGTTTAAGCAGGATTGGAGTTTAATTGGGTGAGGTATTCTCACTGTG
      ACTAAGTTTGATGAATTGAAAAGCGTAGTTGTAGAAAGGAAACTCAAGAAGGAAATTCTT
      GGGGAAACTTAAAGAATCGTATATATGCAATGTCACTTTTTAAGACAACTAATATTTTTA
      AGAATTTACTACTTTTGAGGTGCTGTACTAATATATTACATGTATAATTTCATATATCTT
      [C,T]
      AACTACTAGTTCCTGTAAATAAGTATGCTGATGATGACACGTTCCATTTCTTTCGATAGC
      CACAAAAACAGGAAGTGATGACAAAGCTGGATTCTAACTCCTGACTCCCAAATTCTCTAA
      GACCCTCAGCATTAACATATATTTTATTTTAATGTTATTATATATGTATCATTACTTTTA
      CAACTCTTAAACCAAACATTTTAAAATTAGCTACAACTGCAAAATCAACTTAAAAATTTC
      AAAGAGCCATTTAACATGATAAATTAAAATATTTTAGTAAAACAAAATCACCACTGATAC
```

71994
```
      TTGGAGTTTAATTGGGTGAGGTATTCTCACTGTGACTAAGTTTGATGAATTGAAAAGCGT
      AGTTGTAGAAAGGAAACTCAAGAAGGAAATTCTTGGGGAAACTTAAAGAATCGTATATAT
      GCAATGTCACTTTTTAAGACAACTAATATTTTTAAGAATTTACTACTTTTGAGGTGCTGT
      ACTAATATATTACATGTATAATTTCATATATCTTCAACTACTAGTTCCTGTAAATAAGTA
```

FIGURE 3UUU

TGCTGATGATGACACGTTCCATTTCTTTCGATAGCCACAAAAACAGGAAGTGATGACAAA
[G,A]
CTGGATTCTAACTCCTGACTCCCAAATTCTCTAAGACCCTCAGCATTAACATATATTTTA
TTTTAATGTTATTATATATGTATCATTACTTTTACAACTCTTAAACCAAACATTTTAAAA
TTAGCTACAACTGCAAAATCAACTTAAAAATTTCAAAGAGCCATTTAACATGATAAATTA
AAATATTTTAGTAAAACAAAATCACCACTGATACTTTAATATTCTTAGGTCTGAGAAAAA
CCATTATGTCGTATTATTCCTGCGTTCCTGGTAGCGTTTCTACTGCTGGACATCAGAAAT

72010   TGAGGTATTCTCACTGTGACTAAGTTTGATGAATTGAAAAGCGTAGTTGTAGAAAGGAAA
CTCAAGAAGGAAATTCTTGGGGAAACTTAAAGAATCGTATATATGCAATGTCACTTTTTA
AGACAACTAATATTTTTAAGAATTTACTACTTTTGAGGTGCTGTACTAATATATTACATG
TATAATTTCATATATCTTCAACTACTAGTTCCTGTAAATAAGTATGCTGATGATGACACG
TTCCATTTCTTTCGATAGCCACAAAAACAGGAAGTGATGACAAAGCTGGATTCTAACTCC
[T,C]
GACTCCCAAATTCTCTAAGACCCTCAGCATTAACATATATTTTATTTTAATGTTATTATA
TATGTATCATTACTTTTACAACTCTTAAACCAAACATTTTAAAATTAGCTACAACTGCAA
AATCAACTTAAAAATTTCAAAGAGCCATTTAACATGATAAATTAAAATATTTTAGTAAAA
CAAAATCACCACTGATACTTTAATATTCTTAGGTCTGAGAAAAACCATTATGTCGTATTA
TTCCTGCGTTCCTGGTAGCGTTTCTACTGCTGGACATCAGAAATAGAGAATAGTAGAGCC

72612   CTGAGATAAGAGCAGAGACAGGGGAAAAGCAAAACATTTCTGAAGAGGCAGTTGGTCTAG
TTTGGCTATAATCACTAGACGGGTAAAGGAACATTGGGTGCATTAAAAGTAGAGAGCCTG
GGATGAAGGCGTGAAGGCTGAGTAAGAATCTCTTCACTTGGTAGTAATTCTAGTTCATCC
CCCTCTGACCTGCAATTCTGAACATGGTGTAGCTTGGTCAATAAGGAAATAAATTGCCTT
TCTGGCTGGAGAGGCAAAGGGTAGACAATACATTGTGCCAGCTGAACTTCCTGTCTCTCC
[G,A]
CTCTGGAGAAGAGCCAGTCACAATGTATGACTCAGCACGCCGGGCACCTCTCCCACGCCA
GCCAGGCCTGCCCAGCCACTTGCTGAATCACAAGTGGCCATTTCCAATCCCATCAGTGAC
CCAAGCTCTCCAACTTAGACTAGTTTCTCTGTGATCGGTCTATGATTGTCATGGAGCACA
AAAAGTATTAACTTCTAACATTTATTTTTCTTTCCTGGATGCTTGATGAACTTTATAAGC
AAGAGACTGATTTAATTGTTCCTCATTATCATCTGAGCATGCCGTCTTGGCTTGCCCTTT

73294   TCTGTAGAAAATAATAATAGCAAAATTTCTCCCTTGAGAAGCTTCATAAATTAAATCTCC
AGAGCCAGTATATGTAAGCCGACAGATTATGAAATATGATTTAATGCTCTGTCCAGAGAA
AGGTCAGGGCTTCAGAAAAATCATCATAATATCAAGAAAAACTAATCTGCAACCTGTTAT
ATGATTTTTAAAAATCACCCCCCATCTTTTTTACTGTGCAAACTGTAGATTTTTGTTTAT
TTTATTTGAGGCTATAGTTTATGTCTTGAATCACACACATATGAGTATTACTTTCTGTGA
[A,G]
GTTTTCATGACCCCTGCAATCAAACTTGGGTCCTTCTGTTAGTTTCTATCACAGTATCCT
TCACTTTTCTTTCACAATTCTTGCCATATTCTATAACTACATATTTGTTTGTTAAATATT
TGTTTATCTTTTATAGATGATTGGCTTCAGGAAGAGGGAAACCATGTCCTTTTGTTCAGT
CCTTTATTCTCAGCACCTTGCACAACATGAATATACAAAAAATATTTGTAAAATGACCAT
CGAATGAACAAGTGCTCATTAAGTACCAAGCTATATGCCAGGGGTTGCTGATGGTTAGAA

73385   AAATATGATTTAATGCTCTGTCCAGAGAAAGGTCAGGGCTTCAGAAAAATCATCATAATA
TCAAGAAAAACTAATCTGCAACCTGTTATATGATTTTTAAAAATCACCCCCCATCTTTTT
TACTGTGCAAACTGTAGATTTTTGTTTATTTTATTTGAGGCTATAGTTTATGTCTTGAAT
CACACACATATGAGTATTACTTTCTGTGAAGTTTTCATGACCCCTGCAATCAAACTTGGG
TCCTTCTGTTAGTTTCTATCACAGTATCCTTCACTTTTCTTTCACAATTCTTGCCATATT
[C,G]
TATAACTACATATTTGTTTGTTAAATATTTGTTTATCTTTTATAGATGATTGGCTTCAGG
AAGAGGGAAACCATGTCCTTTTGTTCAGTCCTTTATTCTCAGCACCTTGCACAACATGAA
TATACAAAAAATATTTGTAAAATGACCATCGAATGAACAAGTGCTCATTAAGTACCAAGC
TATATGCCAGGGGTTGCTGATGGTTAGAAATGAGCAGGGCACAAAATTCTTTGTTCAATT
AGTGAGCAATTCAGGCAAAAAGAAAATATTAATGGTGATTATACAATATAATGCAATGCA

74121   CTCTCACCTCAACCTCTTAAGTAGCTGGGACTACAGGTGCATGCCACTATACTGGCTAAT
TTAAAAACAGAAGCCAACAAACAAAAAACACACCTTTTTAAGACTGGGTCTCACTATGTT
GCCCAGGCTGGCCTTGAACTCCTGGCCTCAAGCGATCATCCTGCCTTCCAAAGTGCTACC
TTCTAGAGTATTGGGATTACAAGCGTGAGTCATCTGCACCAGGCCTGAAGCATTCTGTAA
TGGAGAAATACCTGGGTGCTATGGAAGGGCAGAGGGGGAAACACAGAGGAGTAACATCTA
[G,A]

FIGURE 3VVV

TTTACGTTTGTCAAGGAGAGGCCAGGAAAGACTAACTACAGGGGAGATAAACTCCAACCA
AGAGTCTTTAAGTCTTCCAAGACTTACGTACAAGTTTCTTATTGCTAAAATGGAAGTTTT
AATGAACATTTATTTATTTATTTGAGATGGGGTTTCACTCTTGTTGCCCAGGCTGGTGTG
CAATGGCACAATCTTGGCTTACTGCAACCTCTGCCCCCCAGGTTCAGGTGATTATCCTGC
CTCAGCCTCCAAAGTAGCTGGAATACAGGAGCCTGCCACCATGCCCAGCTAATTTTTTTT

75646   CACTCCAGCCTGTGCTACAGAGAGAGACTCCGTCTTAAAAAAACAAAACAAAATAACAACA
ACAACAAACAAAGATAGATGCATAGAGTTTTTCACTGTTGCACTATTTATATTAGCCAAA
AACCGGGAAACAACCTGAATATTCATCAAGTGGGGACAGGTTGAGTAATCATGTGACATA
CATAAATTGCAGCACTGCACACTTGAGAAAAGAAGTGAGAAATGTCTCTATTTCCTAGTG
TGGTTTGCTCTCCAGAGTATACTGTTAAGTGAAAAAAGCACTGTGGCCTCAAATTTATCT
[A,G]
TAGATTCTATACAATCCCCATCAAAATCTCAGCTGGCTTCTTTGCAGAAATTCACAAGCT
GATCTTAAAATGTGTATAGAAATCCAAGGGACTCAAAATTCAATAAATTCAAAGACTAGC
CAAAACAATCTTGAAAAAGAAGAGCAAAGTTGGAGGGCTCATACTTTTCAGTTTCGAAAG
TTGTTATGAAGCTACAATAATCAAGATAGGGTGGTCCTGGCATAAGGATAAACATGGAAC
AGAATTGAGCATCTAAAAATAAAGCCTCATATTTCCAGTCAATTGACTTTTAACCAGGGT

75698   TAACAACAACAACAAACAAAGATAGATGCATAGAGTTTTTCACTGTTGCACTATTTATAT
TAGCCAAAAACCGGGAAACAACCTGAATATTCATCAAGTGGGGACAGGTTGAGTAATCAT
GTGACATACATAAATTGCAGCACTGCACACTTGAGAAAAGAAGTGAGAAATGTCTCTATT
TCCTAGTGTGGTTTGCTCTCCAGAGTATACTGTTAAGTGAAAAAAGCACTGTGGCCTCAA
ATTTATCTATAGATTCTATACAATCCCCATCAAAATCTCAGCTGGCTTCTTTGCAGAAAT
[C,T]
CACAAGCTGATCTTAAAATGTGTATAGAAATCCAAGGGACTCAAAATTCAATAAATTCAA
AGACTAGCCAAAACAATCTTGAAAAAGAAGAGCAAAGTTGGAGGGCTCATACTTTTCAGT
TTCGAAAGTTGTTATGAAGCTACAATAATCAAGATAGGGTGGTCCTGGCATAAGGATAAA
CATGGAACAGAATTGAGCATCTAAAAATAAAGCCTCATATTTCCAGTCAATTGACTTTTA
ACCAGGGTGCCAAGAAAATTCAATGGGGGAAGAATTTGTCTTTTCAACAACTGGTGCTGG

79007   AATATATTTTTAAGTATATTGATTGCTTTTGGAGGGGTTCTAGGAAACAAACAAATCATTT
TGAAAAGTGGTAAATAAAGGAAAGACTTCAGTTCAAGACCAGTCTGAGCAACATAGTAAG
ACCCCATCTCTACAAAAAAATTAAAATATCAGCTGAGCATTGTGGTGTACATCTTTAGTCC
TAGCCACTTGAAGGCTGAGGCTGGAGGATTGCCTGAGCCCAGGAGTTCAAGGCTGCAGTG
AACTATGATGGCACCACTGTGGTCCAGCCAGGGTTAAATAGCAAGACCCTGTTTCTGGCG
[-,A]
AAAAAAAAAAAAAAAAAAAAAGGAAGACTTAAACATACCTTTCCTATATGAACTGTGCCT
CGGAGTAACTAAATAATTGATTAAAGCAAGTTTCTCTGTATAAAAGTACTCCAGCTAAAA
CATTAAGGAGAAATGATAGAATTCAAATATCACAACCCCTAAGGAATTTTTGCATCAAGA
CAACAATAATTAATGACTGATAACACCACACACAGAATACAGACTTATTAATTGTATAAC
TCCTGATCAAGTGCATACCACTATCTGTGAAATAGTTTTGCCAAAAAAAAAAAAAAAAAT

80043   ATTGCAGGCGTGAGCCACTGCACCCAGCCCACCCTTGGTTTTTTTTCAACAAAAAATTACT
AGAAATAAAAGAATAATAGTTGGTCAAGGAAGCTGTAGAATAAGAAAGACTGCCACATAC
ATCAATGGCAGTGGGCGGGCTTTGTTTGAATCCAACTCTAGCATGCAAACATTTGATAAA
AATTTCTTTATTTAAAAAGAAAAGTTTACAAAACAATCAGAAAAAATAAAAAAGATTGAG
GATCTCAGGACAACTACTAGCCTAGATAATTTATAAAGATTAGATAACTGACTCATTTTT
[A,G]
TTAGTTTCTTTCCTAATAAGGCAATATGTATTAGATATATCAGAGTAGAAGGAAATATTT
TTCTTACATCTATTTGGCTTTTTAAAATATAAACATATATAAGTAAAAACCAAAATGATTT
ATAATCCCACCATTTATGTAACTATCTTATTTTCAAAAAAAATTATGCAAATACTAGCAT
TTGTGTGCTTTTTTTTCCTTTTGTGTTTGTGTGTTTATATCCTTTTTAAATATATCCTTTT
TATGTACCTAAGCAGCTGTATACTATACTGCATACTATAGTGTGAACTTTGTTCTTTTCC

80499   AAAAAAATTATGCAAATACTAGCATTTGTGTGCTTTTTTTTCCTTTTGTGTTTGTGTGTTT
ATATCCTTTTTAAATATATCCTTTTTATGTACCTAAGCAGCTGTATACTATACTGCATAC
TATAGTGTGAACTTTGTTCTTTTCCTTCGTCTTTACAACATATTGTGGAAAACGTTCCAT
ATCAGAATATAGATATGCCTTTTTTGTAGCCATTGAAATGCAAAGAAAAAAAGAATATAGA
TCTGTCTCATTTTTTTAAAAATGCTGTATAATCTGTAGCACGAATTTACTATAATTTATTC
[G,C]
CATGCTCCCTTATCGATGGGCATGTAAATTGTGTTAATTTTTATATGATATAATGAGTATC
CTTATATGTATATCTTGGCACAGTTTTTTCGAGTGTATCCATAAAGTTTCTTGCAATGAAA

FIGURE 3WWW

TTATAGGGCAACAAGGGTGTGGTGGCTCTTGTCTGTAATTTCAACACTTTGAGAGGCTAC
GGCAGGAGGATTACTTGAGGCCAGGAGTTTGAGACCAGCGTGGACAACATAGTGAGCCCT
CACCTCTACTAAAAATTAAAAAAAAAAAAAAAGAAAAAGTTTGGTATGGTGATATGTACCT

80940   GGTGGCTCTTGTCTGTAATTTCAACACTTTGAGAGGCTACGGCAGGAGGATTACTTGAGG
CCAGGAGTTTGAGACCAGCGTGGACAACATAGTGAGCCCTCACCTCTACTAAAAATTAAA
AAAAAAAAAAAGAAAAAGTTTGGTATGGTGATATGTACCTGTAGTCCCAGATACCCAGGA
GGCTGAGGTGGGAGGATCATTTGAACCTGGGATGTCAAGGCTACAGTGAGCTATGACTGT
GCCACTGCACTGCAGCCTGGATGACACAGTGAGACCCTGTCTCAAAAAAAAAAAAAAAAA
[A,-,T]
TACAGGCCAAATCCATATGCTTTTAAAGGATATTTTTGAATTGTTCTCAAAAAGAGGCTT
CACCAAATTACCATCCAGGGTATACAAGATACCCATTTCTCCATGTCCTTACCAACAGTG
GCTCTCATCAAGCCTTGGTGGAAATGCTCTCATACTGATACTTTAACGACTAAAAGTCAT
GACATATCTGCTTAGGTTGTAAATTGCCTCCCTCTAAACTTATACAGAGAGAATTTAGAG
TGTTGTCTCAGCTTGGTTCCAGTGTTATCCAAGCCATTAACTTTGTTTTGCCTTAGATT

81615   AGTCTCGCTGTGTCGCCCAGGCTGGAGTGCAGTGGCGTGATCTTGGCTCACTGCAAGCTC
CGCCTCCTGGGTTCAAGCCATTCTCCTGCCTCAGCCTCCCGAGTAGCTGGACCTACGGGT
GCATGCCACCACCCCCGGCTAATTTTTTGTATTTTTAGTAGAGACAGGGTTTCACCATGT
TAGCCAGGATGGTCTCGGTCTCCTGACCTCGTGATCCGCCCGCCTCGGCCTCCCAAAGTG
CTGGGATTACAGGCATGAGCCACTGTGCCTGGCCACAATGGGGTATTGTTTTTATAGACT
[G,T]
TTGAAATCTGCCTTTGGAAACCATGGGTTTGCTGTGTTGTTATGGTGAATGAATTAGGTG
CACAATACTAGTTTTTAAAAAATGAACTTCACACTAGGTACACCTTGAAAAATTATTCCA
GAGCTATAAGAAGAGCTATAAGAAGAAAAATATGATGGGTCATTGCTCCAAAGAAAGGTT
TTAAAATGTAAATTTGTACTTAATGAATAGGACAGTGTACCCTAACCTCCTCCTTGCTAT
TCTTCAGGGATCTCTTCTAACAAGGGCTAATGCTTCACCTAAGCTGTGAAAAGCCTGCTG

82599   CCTTAAAGGAATTTGACTCACAATTTGAAAAACACTGCATTGTAGAATATTTTAGAGTCT
CTTCCCAACCCTCAGAGTCAGATTTATTTCAAGATGGCCCCTGTAAGACAGCTTCAAGCT
TGTGAGTGACTTTCTTTTTTTCTTTTTACTTCTTTACCATTTACCATGACTCCCAAATAAG
TGACTCTTTTGGCTTATTTGGTAACCATGCTAATTTCTACACATAGAACCTAGAGCATTT
ACATAAGACCCACCCAAAGCTTGTGTTTTAACCTTGCTTCTCTCCTTTCTTTCTTTGATT
[C,-]
ATTGATTATGTTTTCTATTGCTATCTGTTCAATCTGTGTTTCAGGCAGTGTACAGGTACT
GAGGCAACAATGGTGAGTAAAAGCAAGCATGCATCCTGAGATATACTGGGAATGAAAGAA
GCTAATCCAAAAGCATACAGGAAAATATTTTTCAAACTTTGATAAATTCTGTGTAAGCATA
TGGCATTGCACGTAACAGGGGAACCGCATTTAATATGGAGTGTTGGAAAAGGCTTCTGTG
AGAAGTGACACTTGAGCTAAGACTAGAAAAGTGAAAAGAATATAACCAGGTACTGGACAG

82952   CAGGTACTGAGGCAACAATGGTGAGTAAAAGCAAGCATGCATCCTGAGATATACTGGGAA
TGAAAGAAGCTAATCCAAAAGCATACAGGAAAATATTTTCAAACTTTGATAAATTCTGTG
TAAGCATATGGCATTGCACGTAACAGGGGAACCGCATTTAATATGGAGTGTTGGAAAAGG
CTTCTGTGAGAAGTGACACTTGAGCTAAGACTAGAAAAGTGAAAAGAATATAACCAGGTA
CTGGACAGCATCATGAGTGCAGGCACAGGTGACATCGTATCACAAGCTTCTAAGGCTGAA
[-,G]
GGGGCGTGAATTGCTAGCTGGAGAGTGGAAGGAAAAGATCTTCAAGATAAAGCTGGAAAA
ATAAACAGGGCCAGGCCTCATAGGTTTCTGTAGACCATGGAAAGAGGTGAAGGTTATTTT
GAGCCTGGATGACATGATAAAACTCACATTGTAAAAATATAACTGCAAGGTAGAGAATGG
ATTGAAGAGGTCCAAGATTACGCAGACAGAGCTATGAACAGCCTATTGCAATGGTCTGGG
TCAAGCATGATGGAGTAGGGTTGGAATAGGGTGGTGAACTTTTATTAGTTATCTTCCTTA

85020   ATCCCTGGGCACTGCTTGCAGGCACTTAATTCTTGATTCAAATGAAACTTTAAAATGTTT
TATCCATGATGTTATGTCTAAAGAAACATGTCAAAGAAACATGTCAGAGAACTTGACTTT
GAATAGAAATCATGGCTGTGCTTTGAGGGAAACAAAATAAATCACAGAGGTAGGAATGCA
TAGTTACAAGCTACTGTTTGTACACAGCAGAGACCAATTCTACTCTCTGTTCTCATTTCC
TCTTCTAATTCCTCATCCCTACACTCCTTCCTGTGTGAAGCCCATGTCTGATCCTGCCTA
[A,T]
TTCAGTGACTGGGGGTCACTGCAGATGCGTGCACAGGGTCCTGTTATGGGATCCGGATTC
TGCCGCCTTCTCCAGACACAAGTTTCCCCTCATACCTGTTGTTCCAGCAAATCCAAGCTA
TTCTCCTTTCCCCACTTGCACTAGGTTCTTTCCCTAGTCTGTGCTTGCATGCATCCTATT
TTTCTCTGGTATTTTTTCAAATTTTTACTTTGGCACCTGGAGAACGTTTTGGCACCACCATT

FIGURE 3XXX

TGTCAGGTGTTTAACTTTGTGCATTTCCTCGTGTGAATGGGAGCGTAGGTCCAGCATCGT

88843   CTTCTGGTGTCAAAAACTATTCAGGTGCATTTCTGTAACCTCTATGCACCTCTCCCCCCA
CCTCCCAGGTGTTATATTTTACAGGCTGTCATACCCTTTTGTACCTCTCCTGAGGAGTTG
TGACATTTGGTGTATAATTAATTCATTTGTCTCCTTTATAAAATTGTGAACTCTGCATGT
TTTGCTTTTCATTGTATAACCAGTATGTGAAAAAAATATGAGCCACATGAATGAATGATT
GACCAGAAGTTCAGGCTTACAAGTAGGAAATATTCAAATATAGGACATTAAATCCAAAGG
[C,T]
CTCAGACCTACTTGTACCTTGGTCTTTACATTAATCATGTTATTTATCATCCAAACCAGG
ATACTCTGAGAGCTAAAGAGGATGCTATTAATATTAATAGCACTGGGAAGAGTCAAAAGC
CATAAATAATCTAGGCAATTCAGGACCTATGTCAACATCATTAAGGCTTTTCAAGGCAGT
GTTTTTTGGTTTTTTTATTTTTTTGTAGAGACAGGGTCTCCCTATGTTGCCTAGGCTGGCCT
TGAACTCCTGGGCTCAAGCAATCCTCCTGCCTCAGCCTCCCAAAACTCTGGGATTACAGG

89700   ATACTGTAATTCTTTTTTAAACCTCCTTCTTCAAAAGAATCAGCCCGATTCATGTTGTAC
TTGAATTCAAGATAACAAAACACCTTTTAGTTACTTAGAAAGATTAGATTGTAAAATATG
TGCTGAGTTCCTAGAAATTAAAAGTGAGAATGAAAAAAAGAATCAATGAAAGTACAGTAG
ATCTCCCGGACAAGGAGAGACCATCTGCATAAAACTGAAGATATAAAATATGTGACTTCC
TACTTTTAGATTAAAAATCTACATTTTGCCTTTGGACATGGTAGAAGATTCAAAATTACCC
[G,A]
TAAACAGTCAGCACTACGTGGAAGTAGGAGCAGCAGTAGGCTGCTGTTTGCTTAGGGTTT
CCTGGGTACCAGGCTGCCTGCTAAGCACTTGTGAGTTATTTCACTCAGTCTTCCCATAGC
TCCAGGAGGTTTATGGCACTTTGTCCCCATTTCACCTTCGATGAAACTCTGGTTCTGAAA
AATTACTTGCCCAAGTTTGCATGGCTATTAAGTAGGGAAAGCATCATGTTTAGGAAATGC
AGAGCTCTTCACCACTCTCCAGCCTGCAGATGCTCAGCATGGCTGCAGCTCTGAGGGGAG

90002   AAACAGTCAGCACTACGTGGAAGTAGGAGCAGCAGTAGGCTGCTGTTTGCTTAGGGTTTC
CTGGGTACCAGGCTGCCTGCTAAGCACTTGTGAGTTATTTCACTCAGTCTTCCCATAGCT
CCAGGAGGTTTATGGCACTTTGTCCCCATTTCACCTTCGATGAAACTCTGGTTCTGAAAA
ATTACTTGCCCAAGTTTGCATGGCTATTAAGTAGGGAAAGCATCATGTTTAGGAAATGCA
GAGCTCTTCACCACTCTCCAGCCTGCAGATGCTCAGCATGGCTGCAGCTCTGAGGGGAGC
[G,A]
CGGGACACCTATGCATGGCCACCTGCCTCAGGCACCCACAGACGAAAGTGGTACATGTGG
AACGGACAGACAGAGAACAGCCTAAAATTGGAAGCTAAATTGTGTGAGAAAGACAAGTAC
TTCAGAGAAGATAGTGTGGAGTCGCAAAATAAGTTTCATGAGAGCTCATACAGAAAACAG
CCTAAAACTAGAAGCTAAATTGCGTAAGAAAGACAAGTACTTCAGAGAAGTTGGTTGGGA
GTAAGAAAGCAAGTCTCATGAGAGCTCTGAGGGGTGTAAATGGGACTTTTAACAGCCAAA

90615   TCTAGCCTAGCAAGAGGAGCTCAATGGATGGAAGTCCTCACTTGTTTCCCTGTGTTAACA
TAGAAGGGGGTCTTTTTAAAATTTTGTTTTTCACTTCAGCTTTTCTGCCAGAAATGTCTAG
TGTAGTGATGTTTTAAAAAAAAACCTAAGTATCTGTTTCCGCCACAAATCCCCATTAAGAC
ATAAATGGAGTTTTATTTTGTGGATGTTTAAAAATCCATGGACTTGAACTTTTGGTAGTT
TCCCAAATATGTAGAATATTCAGCTAGTTTTCTTCAATTTCAGAATCTTTCTTTTCTATC
[A,G]
TTGTTAAAGACACAGGGTTGCATAATAACCATTAAGTTTGAATTGTGCAATTAGACAACT
TTCTTATTAGTCAAGAAGTCAAACTTTTTGTGTGAGTACAGCTTGAAAATCAGCTTTAGT
TTCCAAAGAATGGCCAGTTTGAAGTATAATATTCTCTTTTGCTTACTTGAAATCTGCAAA
TAAATGCTTTAAATTAGGGACAAAGTGATTATTTGCTTTTATTTAAAAAAATAAGGGAAAC
AAAACTCATTACAATCTCTTCTACAGGGTTAGTACTATTCTATTTGTTGATTGCCTCAGC

92506   CAGTCCCAGTGACCAGAAAGAACATACCTTTATCAGCTCGCAGTTTCTTTGGGACAGGTG
TCTGGGCACAGTCTAGTTGAGTTCTCGGCACAGCTGCCATTAAGATGTCAGCCAGAACTG
GGTTCTCTTCTGGAGGCTGAACTGGGCAAGAATCCACTTCCAAGCTCAGTCAGAATGTTG
GCAGGAGGTATTTCCTTGTGGCTGTAGGACCCATGGTGGCTACTTTCTTTAAATTTAACA
AGGAGAAGAATACCGTAGAGTAAGTTGGCTAGAAAGAAAACAGAGTACACATACTTGAAT
[A,G]
ATGATATATAACATTGTAACATAACTCAGTCACAGAAGTAAGACCATCACATCTGCCATG
TAATGTCGGTTAGAAACAAACCATGGAACCAGCCCATGCTGAGGGGCTGGAAATTATGCA
AGGGTGTGAACACCAAAAGCTGGGAATCCTGGGGGGTCACCGTACACAGTCTGTTCACATT
TCCTCTAAAGAAGTTGCACTGCATCACAGTTCCATACCAATTTCTGCTATGACCTTAAAT
ATAGCCCTGAACTTCCCTGTCAAGGAAGAAGTGAGGAGGTTTCAACAAGTGATCAGTAAT

FIGURE 3YYY

92558   GACAGGTGTCTGGGCACAGTCTAGTTGAGTTCTCGGCACAGCTGCCATTAAGATGTCAGC
CAGAACTGGGTTCTCTTCTGGAGGCTGAACTGGGCAAGAATCCACTTCCAAGCTCAGTCA
GAATGTTGGCAGGAGGTATTTCCTTGTGGCTGTAGGACCCATGGTGGCTACTTTCTTTAA
ATTTAACAAGGAGAAGAATACCGTAGAGTAAGTTGGCTAGAAAGAAAACAGAGTACACAT
ACTTGAATGATGATATATAACATTGTAACATAACTCAGTCACAGAAGTAAGACCATCACA
[T,C]
CTGCCATGTAATGTCGGTTAGAAACAAACCATGGAACCAGCCCATGCTGAGGGGCTGGAA
ATTATGCAAGGGTGTGAACACCAAAAGCTGGGAATCCTGGGGGTCACCGTACACAGTCTG
TTCACATTTCCTCTAAAGAAGTTGCACTGCATCACAGTTCCATACCAATTTCTGCTATGA
CCTTAAATATAGCCCTGAACTTCCCTGTCAAGGAAGAAGTGAGGAGGTTTCAACAAGTGA
TCAGTAATGATTCTTTTATGTCTAAGATTCTAGGATGATTTCCTCTCTGCCCTGGTAGGC

92667   AAGCTCAGTCAGAATGTTGGCAGGAGGTATTTCCTTGTGGCTGTAGGACCCATGGTGGCT
ACTTTCTTTAAATTTAACAAGGAGAAGAATACCGTAGAGTAAGTTGGCTAGAAAGAAAAC
AGAGTACACATACTTGAATGATGATATATAACATTGTAACATAACTCAGTCACAGAAGTA
AGACCATCACATCTGCCATGTAATGTCGGTTAGAAACAAACCATGGAACCAGCCCATGCT
GAGGGGCTGGAAATTATGCAAGGGTGTGAACACCAAAAGCTGGGAATCCTGGGGGTCACC
[G,A]
TACACAGTCTGTTCACATTTCCTCTAAAGAAGTTGCACTGCATCACAGTTCCATACCAAT
TTCTGCTATGACCTTAAATATAGCCCTGAACTTCCCTGTCAAGGAAGAAGTGAGGAGGTT
TCAACAAGTGATCAGTAATGATTCTTTTATGTCTAAGATTCTAGGATGATTTCCTCTCTG
CCCTGGTAGGCTGCTCTTCAAAGTATGACCTCCTCATTGTTTCTCTGCTCTACCACACAC
TCATTCCCCTCCAAGAAGGCTGCCCACCTGTAATGACCTGTCTACAGAGCCTGTGATAGT

92803   AATGATGATATATAACATTGTAACATAACTCAGTCACAGAAGTAAGACCATCACATCTGC
CATGTAATGTCGGTTAGAAACAAACCATGGAACCAGCCCATGCTGAGGGGCTGGAAATTA
TGCAAGGGTGTGAACACCAAAAGCTGGGAATCCTGGGGGTCACCGTACACAGTCTGTTCA
CATTTCCTCTAAAGAAGTTGCACTGCATCACAGTTCCATACCAATTTCTGCTATGACCTT
AAATATAGCCCTGAACTTCCCTGTCAAGGAAGAAGTGAGGAGGTTTCAACAAGTGATCAG
[A,T]
AATGATTCTTTTATGTCTAAGATTCTAGGATGATTTCCTCTCTGCCCTGGTAGGCTGCTC
TTCAAAGTATGACCTCCTCATTGTTTCTCTGCTCTACCACACACTCATTCCCCTCCAAGA
AGGCTGCCCACCTGTAATGACCTGTCTACAGAGCCTGTGATAGTGACTTGTGATAAATGG
CTATTAGCACATTTACCAATCAAGGTCCTGTTTGCAATTCGGTTGTGGGTCAAAATTATG
TTTGTTTTAACTGAGGTCTTTAGTTTATTTCAGGCAGAGATCTGGGCTGGAGTGTCACCT

95079   TTTGGTAAATCTTCTTTTTCAGTAGACCACAAGCCCTTGCAAATGTTCTCTTTTTCTAACT
CTGGTAGCAGAAGGACCACTTGAGCCTCAAAACAAAACGGCAGTGCAGTAATGAGGGTAT
TAGGTTGATGTGTTCTATTCAGCACCTGCTCCCGAGCTACCGAATAATGAATGAGCATGA
ATTACACATTGTGAAAACAGGAGAATCTGCCTTCTTTGTGTTGTATGCATCAAGCAGTTT
CAAAAGGGCTTTGCAATTGTGTTTCTCACACAAAGCCACCCATTTGTGAAAACCCATGTG
[T,A]
AAAGGCAAAGAGAACTGTCTGTGTACAGGTTAACATTTAACTAGACTGGCAGAGCTTTTA
ATAATTTCTATAAGGTTAATGGCTTCGTTAATATGCAACCTGTGATTTGGTCCAAGTTAA
ATTTTACTTTGCCCAGAATACATTATAATATAAAGCTTAAGCTTTATTCTTTCAGGTTTA
GTCATTTAACACATAATATTGATCAATTATGCATGTTGGACACAGAGCTCTGAATAGAGC
TTTGAAATATAAAACTATGGTTTTAGTCCTCTTAGAGCTATGATGTTTGGTAGGTTAGGT

95089   CTTCTTTTTCAGTAGACCACAAGCCCTTGCAAATGTTCTCTTTTTCTAACTCTGGTAGCAG
AAGGACCACTTGAGCCTCAAAACAAAACGGCAGTGCAGTAATGAGGGTATTAGGTTGATG
TGTTCTATTCAGCACCTGCTCCCGAGCTACCGAATAATGAATGAGCATGAATTACACATT
GTGAAAACAGGAGAATCTGCCTTCTTTGTGTTGTATGCATCAAGCAGTTTCAAAAGGGCT
TTGCAATTGTGTTTCTCACACAAAGCCACCCATTTGTGAAAACCCATGTGTAAAGGCAAA
[G,A]
AGAACTGTCTGTGTACAGGTTAACATTTAACTAGACTGGCAGAGCTTTTAATAATTTCTA
TAAGGTTAATGGCTTCGTTAATATGCAACCTGTGATTTGGTCCAAGTTAAATTTTACTTT
GCCCAGAATACATTATAATATAAAGCTTAAGCTTTATTCTTTCAGGTTTAGTCATTTAAC
ACATAATATTGATCAATTATGCATGTTGGACACAGAGCTCTGAATAGAGCTTTGAAATAT
AAAACTATGGTTTTAGTCCTCTTAGAGCTATGATGTTTGGTAGGTTAGGTGAAGTAGACA

96495   GGAAGCCACACTGCGATTTTTCCAGATAATTGTGAAACAACTACGGGCCATTACAAAACCA
TAGGAAATTAGAAGTGAGGAGTAATTTTGGAGACTGACAAGCTCTACCTTCATCTAAAGGC

FIGURE 3ZZZ

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AGAATTTCTTCTGCAGTCTCCCTAACAAGGAATCGTTATACCTCAGGGATGGGATAGTCA
CTACCACATAAAGTAGTTCATTTTCAGACATGCATAACCTTAGAAAGTTCTTCTCTTGAT
TTACAATTAGCCTCATAGTTCTGTTGCTGCCTATTGGAGTTTTACTACGTGTACAGTCAG
[G,A]
CAGGGCTTCCATTCAGTCACCACCCATTAGTACTGTTGTACTAGTAATTTATGGATGGCG
TCCATTCTTACTGGTCCATGTCCCATTCTGATTTGTGTTTGTGCCATTTTTAAGTGTTTT
GAATATTAACCCTGGTATCAGATAAACATGGAGTCCTGACTTTTTCCATAATCATGAATA
ACAGTGGAATAGTTACATCAGATTTGTGTGCCACTGTGGTCCCATCTATGAAATAGGGAT
AATAATTGTACCTAGTTCATAAGGTTGTTTGAGGATAGTGTGGAATAAAGTATAAAAAGG
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97070
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ATAGTGTGGAATAAAGTATAAAAAGGGCTTAGCCTGGTTTCTCAAATATTGCAATAAATG
AAACTTAGCATCATGATGCTGTCACAATGGTTCAATGATAATTGAAAACATCGATTCATC
ATTTAGCATCCTCAGCTTATCAGTTTCTCACTATCTAGCTCTTCTTACACTGGACACTTC
CTAATTATTCTTTCAATGTTTTCTGGAAGTTAGTTGAATAATTACTGTGCACCAGATACT
ACACAGTAGTCCCCCTTGATGCATGAGGGATACATTCAAGACCCCCAGTGGATACCTGAA
[T,A]
ACGCAGATATTTCCAAACCCATATATACTATGTTTTTTCCCTTTTGTACATACCTATGGT
AAAGTTTGATTCATAGAGTAAGAGATTAACAATAACTAATAATAGAACAATTATAACAAT
ATGCAGAGTAAAAGTATGTGAATGCAGTCCCTCTCTCAAAGCATCTGATTGTACCGTACT
TACCTATTTTTGAACCACAGTTGACTGTGGGTAAAAAGGAAAACTGCAGATAAGGGGGGA
TTACTATACTACGAGTTTTACATGTACCATTTAACTAAATCATTACGACTCTATAAAGTA
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99913
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GGGTGGGGATTGGTTGTTTTGCTTTGTTTTGTTTTCTCTTCTCTTCTTAGGGGAAAAAGA
CATGCAGGGCTTAGTATTCCAACAATTTGAGAAACCAGGGGGCTGGGATTCATTCATTTT
TATGACAAATAGTTACTCGAGCACCTACTTTATTCTTGGGTACTTTTATGAGTCCAGGGG
CTGCTGCATTGAACAATACAGAAAAGAAGTCCTTTCACTTAGAACTTACGTCCTAGTGGG
GGTTGGGGGTTGGGGGTTGAGAGAATGAAGCATTCTTACAAAGAATGTTAAAAGCGAACT
[A,C]
TGGGCAGGAATTGAGGATATGAGTTTTGATGTATAAAGAAAAAGTGACAAGGTCAATAAT
TGGTGGTCTTAGTGTGATAGATATGCCAGTTTGGAAATTGTATTGAATAAATGCTAGTCA
GGGGCTAGGCTGTAGTTATGAAAAGGAGATGATTAAGGAAGTGAGAATAAGGAAACTATT
GGTGTGGGACGGATGAAAAGATTATTGGAGGCAAGTCAAGGAACTGAGAGGCCAGGGTGT
TAGATGGAGCATTCATGTAGACACTGAAGTCACCAAGAATAATAAATAACAAGTAAGAGG
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102375
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TTAGATGAGACATTGGACTGTGGACTTTTGAGTTATTGCTGAAATGAGTTAAGACTTTGG
GGAATTCCCAGAACTGAGGGTTCCTCCCCATTGTAGACCATATAGGTAGCTTCCAGACGT
TGCCAAGGCATTTGTAAACTGTCATGGTGCTAGTGAGAGTGTCTTTTAGCATGCTCATGT
ATTATAATTAGTGTATAATGAGCAGTGAGGATGACCAGAGATCACTTTTGTCACCATCTT
GGTTTTGGCCAGCTTCTTCACTGCATCTTATTTCTATCAGTGGGGTCTTTGTGACCTGTA
[C,T]
CTTGCAAAAACAGTCCTGCTGATTACTAAATTCCTATCTCACCTATTCAAGATGGAGTCA
CTCTGGTCTGAATGCCCCTGATAAGAGAATCCACAGTGTTCAATTCTCCCCAGTTGATTC
TGAAGCATATCCAGGTTTATTAGCCACTAAGTAAAAATATATTATAGACTACTGTCAATG
AAAGAAACATTTTGTAAGTTATTTCATATTTATTTTTACTTGAGAAGACTGAAAAGGTAA
AGAAGTGATGCTAAAATTTAGAACTAGAAAATCTCAACTTGCTCTAGTAGGAATTTTAAT
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102686
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CAGTCCTGCTGATTACTAAATTCCTATCTCACCTATTCAAGATGGAGTCACTCTGGTCTG
AATGCCCCTGATAAGAGAATCCACAGTGTTCAATTCTCCCCAGTTGATTCTGAAGCATAT
CCAGGTTTATTAGCCACTAAGTAAAAATATATTATAGACTACTGTCAATGAAAGAAACAT
TTTGTAAGTTATTTCATATTTATTTTTACTTGAGAAGACTGAAAAGGTAAAGAAGTGATG
CTAAAATTTAGAACTAGAAAATCTCAACTTGCTCTAGTAGGAATTTTAATAGAGCACACT
[-,A]
AGTTTCTTTTCATTTTCTCTCTCCTGGTATGTGAATAAACAACCTTCCATACTGCAATTT
ACCCTGTAGTGAATTAGATGTTACCCTATTATATTTTGGAGAAACTATATAGTTAGAATC
TAAGCTTAGATAACTTATTTTTATGTTTACAAATCCACTTTCTCTTATACATTTTTCTTA
AATTTTTCTCATATTCTTTCTCTGAATTTGTGGTAAAAATACCCCTTTCCCATTCTATGT
CATGGTTCTTTACGAAGCTTTCTCATCCTCTCCATCCCGAGGGAACTATGTCTCATTTAT
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102687
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AGTCCTGCTGATTACTAAATTCCTATCTCACCTATTCAAGATGGAGTCACTCTGGTCTGA
ATGCCCCTGATAAGAGAATCCACAGTGTTCAATTCTCCCCAGTTGATTCTGAAGCATATC
CAGGTTTATTAGCCACTAAGTAAAAATATATTATAGACTACTGTCAATGAAAGAAACATT
TTGTAAGTTATTTCATATTTATTTTTACTTGAGAAGACTGAAAAGGTAAAGAAGTGATGC
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FIGURE 3AAAA

TAAAATTTAGAACTAGAAAATCTCAACTTGCTCTAGTAGGAATTTTAATAGAGCACACTA
[A,C]
GTTTCTTTTCATTTTCTCTCTCCTGGTATGTGAATAAACAACCTTCCATACTGCAATTTA
CCCTGTAGTGAATTAGATGTTACCCTATTATATTTTGGAGAAACTATATAGTTAGAATCT
AAGCTTAGATAACTTATTTTTATGTTTACAAATCCACTTTCTCTTATACATTTTTCTTAA
ATTTTTCTCATATTCTTTCTCTGAATTTGTGGTAAAAATACCCCTTTCCCATTCTATGTC
ATGGTTCTTTACGAAGCTTTCTCATCCTCTCCATCCCGAGGGAACTATGTCTCATTTATC

102939   CTAGAAAATCTCAACTTGCTCTAGTAGGAATTTTAATAGAGCACACTAAGTTTCTTTTCA
          TTTTCTCTCTCCTGGTATGTGAATAAACAACCTTCCATACTGCAATTTACCCTGTAGTGA
          ATTAGATGTTACCCTATTATATTTTGGAGAAACTATATAGTTAGAATCTAAGCTTAGATA
          ACTTATTTTTATGTTTACAAATCCACTTTCTCTTATACATTTTTCTTAAATTTTTCTCAT
          ATTCTTTCTCTGAATTTGTGGTAAAAATACCCCTTTCCCATTCTATGTCATGGTTCTTTA
          [C,T]
          GAAGCTTTCTCATCCTCTCCATCCCGAGGGAACTATGTCTCATTTATCTTTAGGTTTTCT
          GTATCTTACTACAGTGACTTACCAGAGTAGGTAAATATCTGATGAATAAATGAATACAAG
          ATTTAATTAAGAAGTAATCACATTAAACTAATTGTTCCCTCTCTGATCTCTGTAATATTA
          AGTTTCAAAGTAGTTTCTGGGAAAAGTAGTTAACACAATGATGTATGGATTCAATAAATA
          AGAAAAATGGTGCTCAGGGATTTAACAGAAAGCTCATAAAATGTCAAATCCACAGCAATT

106162   GTTTTATGCGGCTTTGTCTATGCTGGCACATAACTAGTATGTACCAATGTATCTCAGAAA
          AGATATCAAGTTTTCTGTTTAAAAATTTCAGTTTGAGAAAAATCAGTTAAAGAAAAACAT
          AAAAAAGATAAAAGTATATGTGTTATCTAGATTTGTGATATAGGGATATGGCAATAATCA
          AGATGGTGATAAGTGAATGCTGAATTTCAAGAACTACTGATTACACCCTCTAGAATAAGC
          TTTTGCCCGTGATGATTAAATGTGTACGATTTCTTCCTAATATTTATTTTTGTGTATATT
          [G,T]
          GGATTTATTAGAATATCAGGGAAGATCTGCAGGGCACAAAAACTGTATGTTATAAATGTT
          AACAGTGTCAATAAGATCTTTGTTATGTCTTTAGAAGGCTGCTAGATGAGGAGAGTCCTA
          GATCTTAAAGGCTCCTTATTCAATTTTTACAAAAAGGATTTGCAAGTGGAACTGAAACTC
          CAAGTACCATCTATTGCTCATTATTTATTTACCTATTTTTGAGCCTGATTTTCCTGATCC
          CACCTGTGCTCAGGGGGCTAAGAAACACTGGTAATGACCTCTAATTTCAAAGCTCACTGT

106378   CTGATTACACCCTCTAGAATAAGCTTTTGCCCGTGATGATTAAATGTGTACGATTTCTTC
          CTAATATTTATTTTTGTGTATATTGGGATTTATTAGAATATCAGGGAAGATCTGCAGGGC
          ACAAAAACTGTATGTTATAAATGTTAACAGTGTCAATAAGATCTTTGTTATGTCTTTAGA
          AGGCTGCTAGATGAGGAGAGTCCTAGATCTTAAAGGCTCCTTATTCAATTTTTACAAAAA
          GGATTTGCAAGTGGAACTGAAACTCCAAGTACCATCTATTGCTCATTATTTATTTACCTA
          [T,G]
          TTTTGAGCCTGATTTTCCTGATCCCACCTGTGCTCAGGGGGCTAAGAAACACTGGTAATG
          ACCTCTAATTTCAAAGCTCACTGTCATTACTTATTTATGGACTGTCCAAAAAGATTTTTT
          CCACTTTCTTCCAATGCCTTATTTCTTCCTTACCTTTACTGCTTCTGACATTTGAAAACA
          GGGTCTCTGATTCTCAGAAATGTGAGCAATGGTGAGATTTAGCATGAAGGTGACTTTCTT
          TAAAATACCAGCTATCCAGAGCTAGGTACAGTGGCAGGCACCTGTAGTATCAGCTACTTG

107310   AGCTTTATCCTCCCTTCAGAGAACAGTGTTTTCATCCCAGGTCTCATCCATGGCTTCACC
          CTACTTCTATCATTAAGGCATCCTATTCTCCTTCAGTCAACTTCTTCCTCCTCCTCATTT
          TCTTGGTGACTTGGTCATTGCAGATGAGGAAAAACATGAAGAAATCAATTAATCTTCAAG
          TTTAACCACCTTTAGAGACTACCCTTGTGAAAGATTAATTGTGTAACAGTGTGGTTAAGA
          ATGTGACTTCTGGAGCCAGATTGCCTTCATTCAAAACACACTTCACTCATTTCCTAGCCC
          [C,T]
          GAGAGCTTTGACAAGTTGCCTAAACTTTGTCTTAGTTTTTCCAGGGATCAAAAGAATACT
          TACTTAGAAAAAAAATCTTACTTACAAAAGAAATCTTACAGGGATCAAAAGAATACTTAA
          TTAGGGTCATTGTAAAGACTGACCTGATACGTGTGAAGTACTTGATGCAATGACTGTCAC
          AAAGAAATCACTCAATAAAAGTCTAATATTAGTACAATTCTTCTGAGGCAGTCATGGCTT
          TCTTTCCTTGGAAAGGAAGCTGGGACTGCTTCATCTTGTTTTATGTTTCTTTGTCTATGC

108663   ACTTTCCATGTGGAGCTCATATTTGAAGACCTCATTTGCCTTCTCCATCTCCATTTATAA
          TATTTCATCCCTGATGGGCTGTCGCTTGGGCCTCATGTGGAAATTGTAGCCACTGTGAAG
          GGTAACCACCTATCTCTCTGGTGCCCCCTATGCGCATCCCTACAAGTGAGCTGTGTATCA
          CACCATGCTGCTTACATTTTTATGCAACACGATTCAGTAACAGGCAGAAACTTTTATTCT
          TACTGACTCATATTCTTTATATTCATCTGAAAAGATTGACATTTAAAGGAGCCAATTGTA
          [C,A]

FIGURE 3BBBB

AATGGGAAATCCACTGTGTGAATATTTCTTGTACATCAGAATTTGCCTTAAAAATGTTTT
TAACTTAGAGCACATCTGTACTGTTCTCCCCAAATGTCCCATTTACTAGTTCAGAGCAAG
ATGACATTAGGTCTTGGGTGACTCCTGACCCACTATCCTAATGTATATTTTCATTTCCTA
CCAATGTAAGTACCCCATCCAATTCTATCAATACCATAGTGTCTAAAATTCTTGTATTTT
TCTTATTCAGGAAATGCTACAACCAGAGGAACAGTAATGTCTGCCTGACATATCAGAGAA

108876   TCAGTAACAGGCAGAAACTTTTATTCTTACTGACTCATATTCTTTATATTCATCTGAAAA
GATTGACATTTAAAGGAGCCAATTGTACAATGGGAAATCCACTGTGTGAATATTTCTTGT
ACATCAGAATTTGCCTTAAAAATGTTTTTAACTTAGAGCACATCTGTACTGTTCTCCCCA
AATGTCCCATTTACTAGTTCAGAGCAAGATGACATTAGGTCTTGGGTGACTCCTGACCCA
CTATCCTAATGTATATTTTCATTTCCTACCAATGTAAGTACCCCATCCAATTCTATCAAT
[A,T]
CCATAGTGTCTAAAATTCTTGTATTTTTCTTATTCAGGAAATGCTACAACCAGAGGAACA
GTAATGTCTGCCTGACATATCAGAGAAAATGACAATTATGTCATCATCTGTCACTTAGGT
TTCTTAATACCATCCTGTTACAAGGAATAGAGGCAAAAACTCAGCGTAGGAGGTGAGAAA
AAACTGAGGCTGCCATCTTAACAGCCTTTTCATTGCAGAGTCTCAAAATGTACCAAAAGA
TGAAGTGGACAGTGTCCTTTTAAAACAACATACAGTGTAGAATACAGTAACTTATCCCCA

110733   CAACTGACGTGTAATGAGTACTCACCAGAGTTGAGATGTTCTGCTAAGCCAGGCCCTCTT
TTAAAAATGTAATCTCAAACTTTATTAGGTCTCATAATCACCTGGAAGGCTTATTTAAAT
ATTGGCGCCCAACCCACAGAGTTTCTGATTTGTTATAATAGAGTTGAGGGGGGACGGGGC
GTAAGAATCTGCATATCTAACAAGTTCCCAGGTGATGCTGATGCTGCTGATCTGGGCACT
ACATTGTAGGAATCAATTGGCTCTAAAACCTTCTCTACCTTCCACTTCTACATGAGCATA
[C,G]
ATAATCTTGTAGCTGAGTCAGCTTGGAAATCTATGCAGACTAAAGTAGACAGTTGCATGT
CTGGCTGCTCATCTGAATCACCTGTGGAATTTGTTGTTTTTAATACAGATACCTGGCTCT
CCTACAAGTCCCACTGAATTGGAGTTTCAGGAGACCGAAGCCCAGGCACATGTATTTTGC
AAAACTACACTGAAGTTTCTGATAATGACGGATATCAACAATTAAACGCTTACTTCTTGC
CAAATGCTGTGCTAAGTCTCCTGTAATCATTCTTTCATTTAATATTTCTAATAACCTCTT

111546   AGAAAAAGAAAAAAACAGAGACAACCTACGCTATGATAAAGTTATTGAAATCAGGCATTGG
TGCCACTCCAGCAAGAATGAGTGGCTACCTTTTTTTTTTAGATGAGTGCTACCTTTACTTTA
CTGAAATATCATGACATAAACAAAGCCAAAACACTTTCTGCACAAAATAAAATCCTGGTG
ATAAAGGCAGTGGGATTTATGCTTAGCAGCAGGCTGGATACTATCAGGGAGCAGACAAAG
AAGTTTGATACAGGGCTTGTGGACTGTGGGCCCTGGAAGAATCTGATGACATGCCCTCCA
[A,G]
TTACAGCTGTATCTCATCAAAACCACAGACACATGTAAATGGAAATGCCAACACTTCAAG
ATTCTCTGAAAGCAGTTGACTGTCATGCCAACAGCTAACATAATAGGCTTGTTTGCCTGA
GCTTTTGGCACGGCCCTTTTGTTCCCTTTAGCTGTAAATGCAGGGACCCTAGAGCACCTC
ATAGAGTGTGTTCCCTGCCACGTATAAGTATTAGACCCACACTATATTGCTTTGAGTGTT
AAAGCTGAAAGAGACCCTAGAGATCATTTAGTCTACTCCTTCTTTTTTTATGTGAAGGAA

116728   ATTCTCCAGGTTTCCTGGGTTTCCATATTAAGGGCTATTTTCTTGGAACCAAATCAGAAA
ATGTGCATCTGGGTTTCCAGGGTTGGTTTCCATGGTGAGAGAAGTACGGGGAGGCCACCT
TTCTTTCCTCTCCCCAGTGGTTTTAAGTACAATATCTGTATAATGTAATTTTTTCAAAGT
TGGCATTTCTAGTCTTCTCACAAGATAGAACTGGGAAATTGGAACCTAGGAAAAATTCTG
TGCACCTTCCACTTTTACCCTTGTAATTAACAATGACTAATATTTCTTGAAATCTTTCCC
[T,C]
GGACCAGACAAGGTGTTAAATGTTTTACATTCATTTATTTGTTTATTTTTCTCAGCAGCC
CCATGGGGTGGACTATACTTATCACTACTTTATAATGAGAAAAATCAGAAGCTAAATAAT
TTGGCCGAGATCACATGGCTAATAATTGAAAAGTCTAGATTTAAATCAAGCTCTGTCTGA
TTTCAGAAATCAAGCTTTTTCTTAAAAGGAAGATTAATGAGAAATAAAAATATATATTTG
TAAATATTTTTATCTGTGGTTTTTAAATGGTTCTAAGTCAACTTAGTTAGGCTAACATAT

118403   GTAACCTAAATTTTCATTTTATAATGTAACAATAATGATAGCATCATATAGTGAACATTT
ATTGTTCCAAGCACTTTGCTAAGTTTTTAACATTTATTATTAAACTCTCAACCCCATAAA
ATAGGTTTTACTATTGTTTAGATTTTTACAAGTTAAAAAAAAAATCAGGCCCAGAGAGAGAG
AAAGTGATGTGTTCATAATCACACAGCCAGTGATTGGCAGAGCATGAAATTAAACCCAAG
TCTAGAAACATGCCGTGCCTGAGACATGGACGATGATGTGACAATGATGAAGGTAGAATG
[G,T]
CTGACATTGCTAAGCTCTTCCTAAATGTTAAGCACTGTTGTAACTGCATGCATTGTCATT
TAAACTAAAAACAGTTCTGTGAGGCCACTACTATCGTTACAGTTTTATTATTGCATAATA

FIGURE 3CCCC

TATTAACATATAATTAATGTAGTATATTGTATATATAGTACTATTGTTATAGTATATATT
GTTCTCACTTCAGAAATTAGCAGACTGAAAGGTTAAGAAACTTGTTGACTGTGAAGCTGG
AGACAGTCATAGGGGTCTGATGCCAGAGCCCTAACTCTTAACATGCTGCAGTACTGTCCC

118491   AACATTTATTATTAAACTCTCAACCCCATAAAATAGGTTTTACTATTGTTTAGATTTTAC
AAGTTAAAAAAAAAATCAGGCCCAGAGAGAGAGAAAGTGATGTGTTCATAATCACACAGCC
AGTGATTGGCAGAGCATGAAATTAAACCCAAGTCTAGAAACATGCCGTGCCTGAGACATG
GACGATGATGTGACAATGATGAAGGTAGAATGTCTGACATTGCTAAGCTCTTCCTAAATG
TTAAGCACTGTTGTAACTGCATGCATTGTCATTTAAACTAAAAACAGTTCTGTGAGGCCA
[C,G]
TACTATCGTTACAGTTTTATTATTGCATAATATATTAACATATAATTAATGTAGTATATT
GTATATATAGTACTATTGTTATAGTATATATTGTTCTCACTTCAGAAATTAGCAGACTGA
AAGGTTAAGAAACTTGTTGACTGTGAAGCTGGAGACAGTCATAGGGGTCTGATGCCAGAG
CCCTAACTCTTAACATGCTGCAGTACTGTCCCTTTGTTCATGTCAATAAACATGCCTCTG
CTAAAATAGAAACCCACTTCTCTTAATCAATTTTTTATTGTTGAATGTTAGGTTGTTTCT

118888   TCACTTCAGAAATTAGCAGACTGAAAGGTTAAGAAACTTGTTGACTGTGAAGCTGGAGAC
AGTCATAGGGGTCTGATGCCAGAGCCCTAACTCTTAACATGCTGCAGTACTGTCCCTTTG
TTCATGTCAATAAACATGCCTCTGCTAAAATAGAAACCCACTTCTCTTAATCAATTTTTT
ATTGTTGAATGTTAGGTTGTTTCTCATTTTGAAATACAGATAGAGCATCCCAAATCCAAA
ATGCTCCAAAATCCAAAACATTTTGAACACCAACATGACACTCAAAGGAAATGCTCATTG
[A,G]
AGTATTTTGGATTGATTTGGGGATTTGGGATGGCCAACCAGTATAGTGCAAATATTTCAA
AATCTGAAAAAAAAAAATTGAAATGCAGAACACTTCTGGTCCCAAGTATTTCAAATAGGGG
ATACTCAACCTGTACATTTAAATTTGTAGTAAAAATCCTGTTAGCAGAATTATGTCCTGG
AACTTAGTTATTTCTTTGTGATAAATTTTCATTCAATAATAATAGTGTATTCTCTTACTG
AAAATCACTCAAAGAAAATTTTGTGTTCTCACCACAGAAAACAGTAATGTGGGTAATGTG

125444   GGAGAATCACTTGAGCCCAGGAGGTAGAGGCTGTAGTGAGCCAAGATCATGCCACTGCAC
TCCAGCCTGAGCTACAGAGTGAGACCCCATCATTAAACAAAACAAAACAAAAAACAAACA
AACAAAAAACAAGCAAGTTATGTGCTTCCAAAATACAATGATACCATAGCTGTGGGATAG
AGAATCCCATTCCAACATTTCAAAAGAGAAATGGGAAAGAAGGAAGGGGCATCAGCTCCT
AAACAAGTCCAGAACATATCAAAGCAAATTCTATTATATCTTAAAACTCGAGAATAATCT
[-,A,T]
CTTTGAGTTGTTGGTTTGCCCTCTAGATCTACACAGGCATGGGAGCAATCACTCTCATGG
CTGGGGATGGGGAGAGGGGACTTGCTTAAGTGGCTCTCTACAAAGGCACTACCCACATGG
CTCTCTGTGAAGGCTCTGTCTACACAGCTCTGTTGAGTGGTGGTCCTGCCCTTCGAAACA
GAGGTGGAGGCAACCCTGCTCCCCAAGCCAGTGCACTCTGGACCTGTAGTGGGAATGGCA
GCCCTGATGATCTGTGAATCGCCCTCATGATCCTTCTTCCTTTTACTTGAAGGATAGCAC

125810   GATGGGGAGAGGGGACTTGCTTAAGTGGCTCTCTACAAAGGCACTACCCACATGGCTCTC
TGTGAAGGCTCTGTCTACACAGCTCTGTTGAGTGGTGGTCCTGCCCTTCGAAACAGAGGT
GGAGGCAACCCTGCTCCCCAAGCCAGTGCACTCTGGACCTGTAGTGGGAATGGCAGCCCT
GATGATCTGTGAATCGCCCTCATGATCCTTCTTCCTTTTACTTGAAGGATAGCACATGTT
CACAGCTGGATAGCATTACGGTCCCAGCCTGTAAAATCCAAGAAGTCTGACAGCCTTTCT
[T,C]
CATAAATTCAAACTGGCAGCATCTGCTAGTATAATCCCATCTTTATTTCTAGCTTCTGTT
GTGATAACTACTTGATTGTTCAGCTACACTCTAGTGTGCTCTTCAGAACAGGCTTGCTCA
TTTTCTGCAATATGGATAGAAATCTTCAATTTCTGGTTGCTTTTTGCTTAATTATTTTTT
CTTCAATTCAAACATTCCCTTTAACATTTTACTATAAGCAGACAGAAGGAACCAAGTTAC
TCCTTCAAAGTTTTGCTTAGAAATCTCCTCGGCTGGCCTGGTGCAGTGGCTCATGCCTAT

126092   AAGTCTGACAGCCTTTCTTCATAAATTCAAACTGGCAGCATCTGCTAGTATAATCCCATC
TTTATTTCTAGCTTCTGTTGTGATAACTACTTGATTGTTCAGCTACACTCTAGTGTGCTC
TTCAGAACAGGCTTGCTCATTTTCTGCAATATGGATAGAAATCTTCAATTTCTGGTTGCT
TTTTGCTTAATTATTTTTTCTTCAATTCAAACATTCCCTTTAACATTTTACTATAAGCAG
ACAGAAGGAACCAAGTTACTCCTTCAAAGTTTTGCTTAGAAATCTCCTCGGCTGGCCTGG
[T,C]
GCAGTGGCTCATGCCTATAATCCCAGCACTTTAGAAGGCTGAGGCGGGCAGATCACCTGA
GGTCAGTAATTCGAGTCCAACCTGATCAACATGGAGAAACCCCATCTGTACTAAAAATAC
AAAATTAGCCGGGCATGGTGGTGGATGCCTGTAATCCCAGCTACTCAGGAGGCTGAGGCA
GGAGAATCACTTGAACCTGGGAGGTAGATGTTGCAGTGAGCTGAGAACACAACATTGTAC

FIGURE 3DDDD

TCCAGCCTGGGCAATGAGAGCGAAACTCCATCTCAAAAAAAAAAAAAAAAAAAAAGAAATC

127506   CATTCTTTCTGCTGCTGTCAATAGCCCTCTTCTTTGGTCCCACAACACACCATCATGATT
TCTGCATTAAAAATGCCATCTCCCAAGTAATTAACCTATTCACAGTAAGAACAGTTGTTA
GAAGTTGGGGTTATTTCATCATGGTCCAATGGCTTTATCTTGCTCAGGAAATCAAAGATG
AGTGTTTCTAAAGCAAAAAAAAGGAGGATCTCACAATTGTATCTGTTTCATTCACTCTGC
AGGGTCCATTTTACACCCAAACATTCATTAGTTCATTGTTTGTACTCCTGCCTTTCCTGA
[G,A]
GAAGTCATTGTAGCACTATTTCTTAAGTATATTCAAATTTGGATAAGTTAGTCAAATTGA
TGTGAAAGGACCACCCTTGTAAGCCAAATGTGTAAGTCCTACATAGGGATATTACCTGTT
TTTATCTCCTGATGGGCTTTTTTTTTTTCAAGTTTCTAAATAAATCCAGTGAACAAGTAG
ATACGCTACTCATGATTATATAGGAAAACAGAGAAGAGAAACATACACTTACTTAAAAGT
AGAAACATATCTGCTCTTTCCCACTTCACCCTTAATTTTTTTCTCCCCAGCCAATTTACT

127878   CACCCTTGTAAGCCAAATGTGTAAGTCCTACATAGGGATATTACCTGTTTTTATCTCCTG
ATGGGCTTTTTTTTTTTCAAGTTTCTAAATAAATCCAGTGAACAAGTAGATACGCTACTC
ATGATTATATAGGAAAACAGAGAAGAGAAACATACACTTACTTAAAAGTAGAAACATATC
TGCTCTTTCCCACTTCACCCTTAATTTTTTTTCTCCCCAGCCAATTTACTCACCTTCTGTG
GCTGTGCTTCTGTGTTAGACCCTTGCTAGCTGCTTCTGGGGTTCAGAGCAATTGTGCTCT
[G,T]
CCCTCATCTTTTATGACACACCTAGCAAAACAGAAGCAGAGGAGCGAGTTGAAACAGACA
AACGACTATCTGTTATTCTTCAAACATGCCTAGGATTGTATTTAACTATCACCTATCTAA
AAGAGGTATTCTCGCCTGCCTGGAAAGAATTTTGCTAAGAAAATTGTTTCTCTTCTTCCC
ATATTATTTTACCTCTATGCTAGTTCCCTGTGATTTGATATGTCAACTTTGACAAATTCA
TTTTTCTAAAGCACAGATATGACCTTTTTTGTTAAGAAAAAGAAACTACTGTTGCTCCCC

139738   GCTGTGGTTTGTTTTTAATTCATCTCTAAGTATATTCTGATATCTCATGTGATTTCTCTT
TTTGACTCTTTTTTTAAGAGTTTGTTGTTTAATTTCCACATTTTTGTGAATTTTCCAGTT
TTCCTTCTGTTATTGATTCCTACCTTCATTCCAATTATTTCAGTCTTTTTAAATTTTTTG
ATACCTGTTTTGTGGTTTCCTTCCATGGTTTCCTTTAACTCTGAGCATATTCAAGACGGT
TGTTTTAAAATCTCACTCTAGAAAGCTCAATGTTTGAGCTTCCTCAGGACAATTTCTATC
[T,C]
GTTGATTTTAAGTCTTTGAATGGCAATATTTTCCTGTTTCTTTGTGTGCCTTGTGATTTT
TTTTCTGTTGCTATTGAAAACTCGACATTTAAATATGATAATGTGGTAACTCTGGAAATC
AGGTTCCTCCTTTCTTCATGGTTTGCTATTTTTTGATTGTTGAAGGCTGTAGTTATCCAT
TGTTTAGCGACTTCTCCAAACAATGTTTGCAGAGATTGTCTGCTTTGTTGTGTCATCACT
GAAGTTTCTGTTACTTTAGCCTGTGCTCAGCTAATGTTTTGACTGAGATTTAACACCAAG

140261   CTTTGTTGTGTCATCACTGAAGTTTCTGTTACTTTAGCCTGTGCTCAGCTAATGTTTTGA
CTGAGATTTAACACCAAGAGCATTTTTTAAGTTGTTTTTCTTTTCTTAATTTAGTGTTCAC
TTGGTTCCAGTAAACCTTTGAGTGCTTTCCGGAGTTTTGACAAAGTTGGTTTTGACAGTA
TCTGCTTGTTTTTTTTGATGTTTCTGTTCAGAGATGGGGCTTGGAACTGCTTACATCAGCA
TTTTTCTCTAGATTCTTCTAATCTTGTACCCCAGGTTCAAAAATAAAAGGTACTTTGCTT
[C,T]
AAAACAAAGAATAGTCTTTCTTCCAAGAAGAATCAGAAAGATTATGAACTATTTTTCTGA
TTCTTCACTCTATTTTCTCTCTTTTACATTAAGGCTTTTAAAACATGAGTCAATCTTACC
TTATTATATTATTAACATGCTCGTTCATTCATTCATTCATTTATTCAGATGACTGTAAAA
TTCCTGCTTTGTTAGGAAATATTTCTGACTAGGTGGTTAATGCTATGGTTAGATACACAA
AGTGCTGTGGGAATTGCTCACTGGACCTGAGTGAAGGGTTAGGATAGGCTTTCCAGAGGA

141590   AATGGGAATAGCATTGAATCTATAAATTACTTTAGGCAGTATGGCCATTTTTATGATATT
GATTCTATCTGGGAACCTGGAATGTTTTTCCATTTGTTTGTGTCCTCTCTGATTTCCTTG
AGCAGTGGTTTGTATTTCTCCTTGAAGAGGTCCTTCATTTCCCTTGTTAGCTATATTCCT
AGGTGTTTTATTGTTTTGTAGCAGTTGTGAATGGGAGTTCATTCATGATTTGTCTCTCTG
CTTGCCTGTTGTTGGTGTATAGGAATGCTAGCAATCTTTGCACATTCATTTTATATCCTG
[T,G]
GTTTCAGTATTTTAAAAACTTACTTCAGGTGATTCTATGTGTGCAACCATGATTGAGATA
CACTGTTATAGAATCTAGGATGTGATAAACTAGAAGAACATAACTAAAGTTTTGCATTTT
TCGGGTGTCTCAGTTTCCTCATTTATAGATGGAGTTGGTATGTGTACCAAGTTCATAGGC
TTGTTCTGAGTAAATTAGTGCATGTAAAGTGCTCCACAGAATGTTAGCTGTTGTGATGCT
TTACTTTCCATTGCACTTCCTGACTCCTAGCCTTTCTTTTCCTTGGCTCTTTTTATGCTC

FIGURE 3EEEE

142613   ACAAGTTGAACATGAACCCTTTAAGGGTAATGGGGTCTGAAGTGTCACACTAAAAGGTCA
          TCTGCAAGTATGTATTTCATATCTTTGTTTAAATAAAATAGTTACATAGTAGAGGGAAAA
          AAAATCCATGTGGATTTTGCATTTCACTCAATTATAACCTTGATTTTTAATGCTAAAAAT
          TATTTTTCCTAAAATCTTGGGGTAAAAGTGTTGCTCCAAAGAGCTTTTATCAGATTATGT
          TTATCCTGTAGCTGCCTGTCCCCTGTGACCGATACTGGAAACCCTCAGGATTACAAATGC
          [C,T]
          TCCGTTTGCAAGTAAGAGTGAAATACAGCAGAACTGTGTCTTCTCCTTTGTCTTGTTCCC
          CATCTCTCTTCTGTGCTTTGTATTGTTTCCTCTCCTGTCACCTAAACAGGCACTCTGAAA
          GAAAACTCTCCAGTACTGGAGAACTTAGCATATTCTAATTCCTAGGTTAAAAAAAAATAA
          TAAATGACTGAATGATTTTTTTTTAAAGAATATTTTCCATCAGAAGAAATTTGGAAGTATT
          TTGTTGCAGAATTTTAAAACATTTGATCTGGGTCTAATTCTGTCCTGGGACTGGTAATCA

142774   GATTTTTAATGCTAAAAATTATTTTTCCTAAAATCTTGGGGTAAAAGTGTTGCTCCAAAG
          AGCTTTTATCAGATTATGTTTATCCTGTAGCTGCCTGTCCCCTGTGACCGATACTGGAAA
          CCCTCAGGATTACAAATGCCTCCGTTTGCAAGTAAGAGTGAAATACAGCAGAACTGTGTC
          TTCTCCTTTGTCTTGTTCCCCATCTCTCTTCTGTGCTTTGTATTGTTTCCTCTCCTGTCA
          CCTAAACAGGCACTCTGAAAGAAAACTCTCCAGTACTGGAGAACTTAGCATATTCTAATT
          [C,A]
          CTAGGTTAAAAAAAAAATAATAAATGACTGAATGATTTTTTTTTAAAGAATATTTTCCATCA
          GAAGAAATTTGGAAGTATTTTGTTGCAGAATTTTAAAACATTTGATCTGGGTCTAATTCT
          GTCCTGGGACTGGTAATCATCTTTTTTTTGAGGCTAAATTTTCTCATTTTGATGAAAAAGT
          CATCAATAGATGTTGAAAGCTGGACAGTGCAGTGTCAAAGCAAATGCTTTGCATGTCTGC
          AAGAAAGTCACAAATAAAGAAGGCTCTGCTGACTAAAAGAGAAAGATACTTAATCAACTC

143288   GTCAAAGCAAATGCTTTGCATGTCTGCAAGAAAGTCACAAATAAAGAAGGCTCTGCTGAC
          TAAAAGAGAAAGATACTTAATCAACTCCAGTACCATTGTTGAGGGGAACATTCTATCAGG
          ATTCAGTATAGAGAGATATTTTTAGGCTATTCACAAAATCCAGGTAGAACCTCCAAGCTA
          CATTTACAATAATACTAGCTTTTAGATTAATTGTTGTTTTTTAAATATGTATTAGCCTCT
          TATACAAATATAAGGAGTTACAAATTATTATTACAATAATCTTGGCTTTCGTGATTGTCC
          [G,A]
          ATGTATTTACACGTACCGAGAGCTTTATTTCTCCGTATAGTTTCAAGTTACTGTCTCGTG
          TCCTTTCATTTCACCTTGCAGGACTCCTTTGAGCATTTCTTACAGGGAAGTTCTAGTGGT
          AATAAACTCCCTCCACTTTTATCTGGAAACATCTTAGTTTCTCTCTCACTTTTCAAGAAC
          AGTTCTGCCAGATAGAGGACCCTTGGTTGATAGGTTTTTTTTCTTTTAGCACTTTGAATAT
          ATCAGCCCACTGCCTTCTGGCCTCCAAAGTTTCTGATAAGAAATCTGCCCGTCATCTTAT

145610   TTTCCACCCTGACTCTCACAGGCTGTGTGCAAACTGCTCCGGAACATGTGTGTGCTCAGC
          TCCCTCCCATGGGGCTGGAGGATGAGGGATGGGTAGCTGCTGCTGTGCTAAGAGCTTAAG
          TTGGTCATAATTAACTGCGCTTTGCCACCCAAGCCTTCCCTGAAAGTTGCAAGCTTTCAA
          TAGACTCCAGAGTTCTAAAATAGTGACATTAGACAGATTCTGCCAGTGCAATCGCTGTCT
          AGGAGGGGAGACAGATTCCTGGTGCTTCCTGTTTTGCCAGCTTCCCGGAATCTTCTTCAC
          [A,C]
          TAGCATCCATTTTGAAGATACTACTTACTTCTCAATTTGGGGCTATTCATTGAATAGACT
          GTCACCAGGTTATTGGCTGTTTGAAGATTCTCATTTGTCTGCTAACTATACCTCTATTTT
          TTTTCTACGTTCACCTGGAAGACATGTCTTCTTCAAGAGCACCTTGACTCTGTCCAGAAG
          GAGTTCATAATTTTCAACAGAGAAAGTAAGTAATTCCTGGGAGAACAACAGCCCCAGAA
          ATGGTGGCATGTTTCAGCCAGACTTTACTTGCAGAGAAAATATATTTTTAACATTTTAAA

148360   GAAGTGAACAGGCTTGGTGGGGGATTGTTTTCACCTCTTGGCTACTCAGAGTACCTAAAC
          CTGTCCTTACTTATGGAGAGCATGTGTCACACCAAGATGGCAGTAAGCTGGCAACTGCGA
          AGACCTGACTGATGCCCATTTGGGAAGCCAGGCAAGTGAAAATGGACCGAAGAAACAGAG
          ATGGCTGTCTTTTATGCAGGGCTTTTCCATAAAGAGGTTACACTGGGGCAACCAAGTATG
          TGTAGAAAGCCAGAGCTAAACTTCAGCTTGGCATTCACAGTTTTCTCTTCACTGAGCTAA
          [T,C]
          AGGCCCAGAGTTTCGGGCAGAGCTGTGAAATAGTGCTTCTCTAATAGCAACCATATTATT
          GTTACATAATTAAAAGCCAGCTCTTTTGTTGTTTGTTTGATTCCTTTTCCCTACAGTTCC
          CACATCATTTGTCTGTGCTATTCTGTTTTTTCTCCAAACACTATAAACTTGAAGCAATTGC
          CCTGACTCGATTTCAGAGAAGGGGATG

Chromosome map:

Chromosome 5

FIGURE 3FFFF